

1 **The Clinical Presentation and Genotype of Protein C Deficiency with Double Mutations of the**  
2 **Protein C Gene**

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4 genetic analysis. YK, TH and TU managed the quality control of the screening of thrombophilic  
5 predisposition. SO and DK organized the nation-wide cohort of pediatric thrombophilia in Japan.

6

7 **Abbreviations list**

PC	protein C
PF	purpura fulminans
PS	protein S
FV Leiden	factor V G1691A
FII variant	prothrombin G20210A
<i>PROC</i>	protein C gene
Rr	reference range
PCR	polymerase chain reaction
Pt(s)	Patient(s)

ICTH	intracranial thrombosis and hemorrhage
DVT	deep vein thrombosis

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1 **Abstract**

2 **Background.** Severe protein C (PC) deficiency is a rare heritable thrombophilia leading to the  
3 thromboembolic events during the neonatal period. It remains unclear how complete PC gene (*PROC*)  
4 defective individuals develop or escape the neonatal stroke or purpura fulminans. **Procedure.** We  
5 studied the onset of the disease and the genotype of 22 PC-deficient patients with double mutations in  
6 *PROC* based on our cohort (n=12) and the previous reports (n=10) in Japan. **Results.** Twenty-two  
7 patients in 20 unrelated families had 4 homozygous and 18 compound heterozygous mutations.  
8 Sixteen newborns presented with purpura fulminans (n=11, 69%), intracranial thromboembolism and  
9 hemorrhaging (n=13, 81%), or both (n=8, 50%), with most showing a plasma PC activity of <10%.  
10 Six others first developed overt thromboembolism when they were over 15 years of age, showing a  
11 median PC activity of 31% (range: 19-52%). Fifteen of the 22 patients (68%) had the 5 major  
12 mutations (G423VfsX82, V339M, R211W, M406I, and F181V) or 2 others (E68K and K193del) that  
13 have been reported in Japan. Three of the 6 late-onset cases, but none of the 16 neonatal cases, had the  
14 K193del mutation, which has been reported to be the most common variant of Chinese thrombophilia.  
15 A novel mutation of A309V was determined in a late-onset family of two patients. **Conclusions.** The  
16 genotype of double-*PROC* mutants might show less diversity than heterozygous mutants in terms of  
17 the timing of the onset of thrombophilia (newborn-onset or late-onset).

18

## 1 **Introduction**

2 The development of thromboembolism depends on the genetic background of patients and triggering  
3 factors. The presence of protein C (PC), protein S (PS) and antithrombin deficiencies is associated with  
4 a high risk of inherited thrombophilia. Patients who are heterozygous for either gene mutation develop  
5 venous thromboembolism earlier than the carriers of factor V G1691A (FV Leiden) or prothrombin  
6 G20210A (FII variant) [1]. In contrast to the high allelic frequency of FV Leiden or FII variant in  
7 Caucasians, neither of these mutations has been found in Asian patients. These natural anti-coagulant  
8 deficiencies are thus the leading genetic cause of Asian thrombophilia [2]. Recent studies [3,4] and a  
9 nationwide survey [5] in Japan have shown that the major inherited thrombophilia in pediatric patients  
10 is protein C (PC) deficiency, while protein S (PS) deficiency mainly affects adult patients due to the  
11 high prevalence of PS-Tokushima K196E [6,7].

12 Severe PC deficiency is an extremely rare thrombophilia due to complete PC defects, presenting  
13 as neonatal purpura fulminans (PF) in both Asian and Caucasian populations. Infants with double  
14 mutations of the PC gene (*PROC*) develop neonatal PF and/or stroke [8,9] and rarely escape the  
15 development of thromboses during infancy and childhood. Fetal distress and infections can predispose  
16 individuals who are heterozygous for *PROC* mutations to neonatal and pediatric thromboses [10-12].  
17 However, little information is available on the onset of disease in PC-deficient individuals who harbor  
18 double mutations of *PROC*.

1 We herein report the first presentation and genotype of patients with biallelic *PROC* mutations  
2 in Japan in our own cohort and in an extensive review of the literature. Notably, six patients with  
3 double mutations of *PROC* developed no overt thromboembolic events until they reached 15 years of  
4 age. The genotypes and phenotypes of patients with neonatal- and late-onset thrombophilia are  
5 discussed.

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## 1 **Patients and Methods**

### 2 *Subjects and data collection*

3 Clinical data were collected from our cohort genetic study and a literature review. Twelve patients were  
4 diagnosed with biallelic *PROC* defects, as assessed by the thrombophilic screening of plasma PC  
5 activity and/or a genetic study at Kyushu University from 1993 to 2015. This study was certified by the  
6 Institutional Review Board of Kyushu University (#232-02) and Yamaguchi University (H26-136).  
7 Written informed consent was obtained from all of the subjects. The collected data included gender, the  
8 age at the onset of each thromboembolic event, the family history of PC deficiency, PC antigen and  
9 activity levels, genetic study results, and the outcomes. We further reviewed all of the publications and  
10 sentinel sources, including meeting reports in Japan, using the Japana Centra Revuo Medicina,  
11 PubMed, and Google Scholar for citations that were published from 1981 to December 2015. The  
12 search terms were congenital, inherited, hereditary or heritable PC deficiency, compound heterozygous,  
13 PF and thrombophilia. We collected 22 PC-deficient patients for our analysis, including 12 from our  
14 cohort and 10 from the extensive review [13-19].

### 16 *Coagulation study*

17 Coagulation tests were performed in our cohort as described previously [4]. The anticoagulant activities  
18 of PC were determined using the Staclot Protein C kit (Diagnostica Stago, Asnieres, France). The



1 reference ranges of the PC activity/antigen levels for term and preterm infants and other subjects were  
2 based on those reported in previous studies [4,6,8,20]. In our cohort study, the plasma PC activity was  
3 determined using repeated coagulation tests (including factor VII activity) at the time of diagnosis and  
4 during the disease course.

5

### 6 *Genetic analyses*

7 Genomic DNA was extracted from peripheral blood leukocytes after obtaining informed consent from  
8 the patients. The direct sequencing of polymerase chain reaction (PCR) products was performed for the  
9 coding regions of *PROC* (exons 1–9), as described previously [6]. The exon and exon-intron boundary  
10 regions of each gene, including the promoter region, were amplified by PCR, and the products were  
11 then subjected to direct sequencing using an ABI 377 (Applied Biosystems, Foster City, CA, USA).

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1 **Results**

2 *Clinical presentation of PC-deficient patients with double mutations*

3 The characteristics of all 22 patients who had double-allele *PROC* mutations are summarized in **Table**  
4 **1**, based on our cohort and a review of the cases that were reported between 1985 and 2015 [4,6,9].

5 There was no declared consanguinity in the 20 unrelated families, including twins (Patients [Pts] 12-1  
6 and 12-2) and a child (Pt 16-1) and his mother (Pt 16-2). They had 4 homozygous and 18 compound  
7 heterozygous mutations. Sixteen newborns first presented with either PF (n=11, 69%), intracranial  
8 thrombosis and hemorrhaging (ICTH) (n=13, 81%), or both (n=8, 50%). Of the 16 neonatal-onset  
9 patients, 2 had prenatal cerebral lesions that were assessed by imaging analyses, and 11 and 6 infants  
10 presented within 7 days and within the first 24 h after birth, respectively.

11 Six patients first presented after the neonatal period. Two developed deep vein thrombosis  
12 (DVT) in adolescence and early adulthood, while 2 others were elderly patients with cerebral  
13 infarctions. Pt 16-1 first presented with epilepsy at 3 years of age but did not experience overt  
14 thrombosis without anti-coagulant therapy until 12 years of age. Pt 17 was diagnosed with PC  
15 deficiency at 22 years of age, when a subclinical hypercoagulable state was identified during the  
16 persistent febrile period of cervicitis (high D-dimer levels and sustained low levels of plasma PC  
17 activity). The detailed clinical information of Pts 16-1, 16-2, and 17 in our cohort are shown in the  
18 **Supplemental Table S1**.

19

1 *Plasma PC activity and the genotype of the neonatal- and late-onset cases*

2 All but one of the neonatal onset patients (Pt 11) showed a plasma PC activity of <10% (the lowest  
3 detectable limit) at the time of their diagnosis. In contrast, the late-onset patients showed a median PC  
4 activity of 31% (range: 19-52%) at the time of their diagnosis. In Japan, 5 recurrent mutations (ex7:  
5 c.541T>G, p.F181V, ex7: c.631C>T, p.R211W [PC-Tochigi], ex9: c.1015G>A, p.V339M, ex9:  
6 c.1218G>A, p.M406I, ex9: c.1268delG, p.G423VfsX82 [PC-Nagoya]) were reportedly found in 49%  
7 of Japanese families with PC deficiency [19]. Fifteen of 22 patients (68%) had these 5 major mutations  
8 (p.G423VfsX82 [n=7], p.V339M [n=4], p.R211W [n=3], p.M406I [n=3], and p.F181V [n=7]) or 2  
9 other reported mutations (p.E68K [n=3], and p.K193del [PC-Tottori] in [n=73] in Japan. PC-Nagoya  
10 was the most frequent allele (22%; 7/32) in the neonatal onset cases (44%; 7 of 16 patients). In  
11 contrast, PC-Tottori was the most frequent allele (25%; 3/12) in the late-onset cases (50%; 3 of 6  
12 patients). Two of the 5 major mutations in the neonatal cases (G423VfsX82, V339M) were not found  
13 in the late-onset cases, while the other 3 mutations in neonatal cases (R211W, M406I, and F181V)  
14 were found in late-onset cases. The novel mutation p.A309V was found in a late-onset family (Pts  
15 16-1 and 16-2).

16

## 1 Discussion

2 The present study first characterized the onset of disease and the genotype of PC deficiency in patients  
3 with double mutations. Early-onset patients presented with PF and/or ICTH within 24 h after birth,  
4 with PC activity levels of <10%. Late-onset patients developed overt DVT at >15 years of age, with a  
5 PC activity of >20%. The genotypes of the neonatal cases were highly restricted to the major Japanese  
6 mutations [6,7,19]. In contrast, the late-onset patients showed genotypes that were distinct from the  
7 neonatal cases, including p.K193del and a novel mutation. The distinct mutation spectrum in patients  
8 of this ethnic background may explain the first presentation of PC-deficient patients with double  
9 mutations.

10 The first survey of double *PROC* mutations demonstrated that there were 2 types of disease: the  
11 majority of cases were classified as neonatal onset (72%), while in the remaining late-onset cases,  
12 thromboembolic events first occurred after adolescence. The present cohort corroborated the finding  
13 that the majority of PC-deficient patients with double *PROC* mutations first presented with neonatal  
14 PF and/or ICTH [9]. More than 200 *PROC* mutations have been reported. The high allele frequency of  
15 PC-Nagoya (44%) in the newborns was as we previously predicted [3,9]. The onset age and mode of  
16 the late-onset patients with double mutations were varied. Late-onset patients with the double  
17 mutations of *PROC* are rarely reported in Caucasians [21,22] but are increasing in number in Asian  
18 countries [18,19,23-32] (**Table 2**). This discrepancy may be due to the high allelic frequency of FV  
19 Leiden and FII variant in Caucasians, which hampers the detection of rare double *PROC* mutations in

1 adult patients. Manabe et al. [32] previously reported a late-onset case in a patient with genetically  
2 undetermined PC deficiency (<5% of PC activity) who first presented with DVT at 14 years of age.  
3 Although he was also found to have dysplasminogenemia, which might augment hypercoagulability,  
4 the late onset of the patient's disease was not explained. Iijima et al. [18] reported a patient with  
5 homozygous K193del as PC-Tottori, who first developed DVT at 23 years of age, with a non-severe  
6 PC activity level of 52% (**Table 1**, Pt 18). Recent large studies in Japanese adults have reported no  
7 patients having double mutations of *PROC* [6,7]. In contrast, Miyata et al. [19] previously recognized  
8 K193del as a polymorphism because the PC activity levels were subnormal, as shown in the 2 elderly  
9 patients with PC activity >60% (**Table 1**, Pt 19 and Pt 20). K193del has been recently identified as the  
10 most common variant in Chinese patients, although the amidolytic PC activity-based screening system  
11 does not discriminate the variant carrier from non-carrier [33]. K193del, M406I and R211W continue  
12 to be found in Korean [31] and Japanese patients with late-onset double *PROC* mutations (**Table 2**). In  
13 this line, the variant allele of K193del may be prevailing as a type II deficiency in Asian patients with  
14 late-onset double *PROC* mutations.

15 A late-onset family (Pt 16-1 and Pt 16-2) harbored a novel mutation of c.926C>T in exon 9. Pts  
16 16-1 and 17 were diagnosed with moderately severe type I and type II PC-deficiency, respectively.  
17 The levels of PC activity in Pt 16-1 were >10% during infancy and >20% during his school life (*data*  
18 *not shown*). A PC activity level of 20% was the hallmark of the late-onset cases (**Table 1**). The disease  
19 onset of biallelic PC deficiency might depend on the genotype that retains 20% of the PC activity  
20 levels in daily life.

1 A major concern is the *true* prevalence of inherited PC deficiency. Pts 16 and 17 were found in  
2 our pediatric thrombophilia cohort [4,9]. The low allelic frequency of *PROC* mutations in patients of  
3 other ethnicities (<0.5%) was originally estimated from a population study based on the assessed  
4 plasma activity. A single assay of natural anticoagulants lacks the power necessary to determine the  
5 inherited deficiency. The asymptomatic parents with a heterozygous *PROC* mutation, such as F181V  
6 and K193del in this study (**Supplemental Table S1**) and our recent report [11], probably show a  
7 normal PC activity level. In pediatric screenings, physiologically low concentrations of natural  
8 coagulants also hamper reaching a diagnosis of inherited thrombophilia. Thus, the *true* frequency of  
9 Japanese carriers of a heterozygous *PROC* mutation may be higher than the previous reports. It is  
10 necessary to clarify the *true* frequency that we would find out clinical and laboratory signs in  
11 asymptomatic variants. Further studies should be directed towards the early diagnosis of  
12 asymptomatic PC deficiency in various conditions, especially for the late-onset type of double  
13 mutations.

14  
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14

1 **Legends**

2 **Supplemental Table S1. The clinical profiles of three patients with the late-onset type PC**  
3 **deficiency in our cohort**

4 Pt, patient; DVT, deep vein thrombosis; PC, protein C; n.a, not available; rr, reference range; *PROC*,  
5 protein C-gene; c.hetero, compound heterozygote; homo, homozygote; PTE, pulmonary  
6 thromboembolism

7 \*: The details of the *PROC* mutation are shown in **Table 1**.

8 **Case presentations**

9 ***Pt 16-1 and Pt 16-2***

10 Pt 16-1 and Pt 16-2 were a 12-year-old boy and his 45-year-old mother (at the time of writing this  
11 manuscript). Pt 16-1 was born at 39 gestational weeks, weighing 3,752 g, by an elective caesarean  
12 delivery to a PC-deficient mother. The family history revealed thrombosis in the mother's but not  
13 father's relatives (Supplementary Figure 1A). None of the family members had received a genetic  
14 diagnosis. The mother (Pt 16-2) suffered from DVT and a pulmonary thromboembolism at 15 and 17  
15 years of age, respectively. Despite the administration of warfarin therapy, she experienced recurrent  
16 abortions because of DVT at 20, 25 and 26 years of age. She gave birth to the boy (Pt 16-1) at 34 years  
17 of age after a pregnancy course that involved prolonged heparinization and the replacement of  
18 plasma-derived activated PC (Anact C®). Pt 16-1 lived an active neonatal life, with a PC activity of  
19 13% at 5 days after birth (Supplementary Figure 1B). Thereafter, the PC activity levels were

1 approximately 20%. Partial epilepsy occurred at 3 years of age but subsided at 5 years of age.  
2 Magnetic resonance imaging and electroencephalography indicated no abnormalities. A physical  
3 examination revealed that the boy had no abnormalities, and he has continued to enjoy participating in  
4 a judo club after school. Coagulation studies revealed a normal prothrombin time and activated partial  
5 thromboplastin time. The plasma levels of protein induced by the absences of vitamin K or  
6 antagonist-II were undetectable. The fibrinogen concentration was below normal (165 mg/dL,  
7 reference range: 200–400 mg/dL), but none of the fibrinogen degradation products or D-dimer levels  
8 increased. The patient's thrombin-antithrombin complex,  $\alpha$ 2-plasmin inhibitor-plasmin complex,  
9 thrombomodulin, plasminogen activity, and plasminogen activator-tissue plasminogen activator  
10 inhibitor-1 complex levels were all normal. The patient's antithrombin (110%) and PS activity (73%)  
11 levels were normal for his age. The plasma PC activity (27%) and antigen (21%) levels were low for  
12 his age (Figure 1B). The patient was negative for lupus-anticoagulants. A genetic analysis using  
13 peripheral blood-derived DNA revealed a reported mutation of c.541T>G, p.F181V in exon 7 and an  
14 unreported one of c.926C>T, p.A309V in exon 9. His 49-year-old father, who had no thrombotic  
15 history, showed a PC activity of 80% (reference range: 75-131%) and carried a heterozygous mutation  
16 in exon 7 (c.541T>G). The exon 9 mutation of c.926C>T, but not the wild genotype, was found in the  
17 mother.

18  
19 **Pt 17**

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1 Pt 17 was a 22-year-old female patient who was referred to us because of a transient increase in her  
2 D-dimer levels during the febrile period of cervicitis. Subclinical coagulopathy subsided uneventfully  
3 without anti-coagulation therapy. She showed no abnormalities on physical and laboratory  
4 examinations. She had no personal or family history of illness. Coagulation studies revealed a normal  
5 prothrombin time and activated partial thromboplastin time. Protein induced by the absence of vitamin  
6 K or antagonist-II was undetectable. The patient's fibrinogen, fibrinogen degradation products,  
7 D-dimer, thrombin-antithrombin complex,  $\alpha$ 2-plasmin inhibitor-plasmin complex, thrombomodulin,  
8 plasminogen, and plasminogen activator-tissue plasminogen activator inhibitor-1 complex levels were  
9 all normal. The patient's plasma PC activity (17%) and antigen levels (50%) but not antithrombin  
10 (99%) or PS activity (78%) were low. The patient was negative for lupus-anticoagulant. A genetic  
11 analysis revealed compound heterozygous *PROC* mutations of c.577-579delAAG, p.K193del in exon  
12 7, and c.631C>T, p.R211W in exon 7. A family study was not performed because informed consent  
13 could not be obtained.

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