



Case report

A new *ENG* mutation in a Japanese family with hereditary hemorrhagic telangiectasia and pulmonary arteriovenous malformations

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ABSTRACT

We present a case series of four siblings with hereditary hemorrhagic telangiectasia (HHT) and pulmonary arteriovenous malformations (PAVM). The patients' mother has HHT. Case 1: A 22-year-old man developed dyspnea and epistaxis. CT revealed a large PAVM, treated by segmentectomy. Case 2: A 27-year-old woman developed epistaxis and dyspnea. CT revealed three PAVMs, treated by partial resection. Case 3: A 20-year-old woman developed dyspnea. CT revealed multiple PAVMs, treated with endovascular occlusion of the largest one. Case 4: A 12-year-old woman developed epistaxis. CT revealed multiple PAVMs, observed without treatment. Genetic testing identified a new mutation, *ENG* c.1517T > C (p.Leu506Pro), in all patients and their mother. We suspect that HHT in these patients may be associated with this *ENG* mutation.

1. Introduction

Hereditary hemorrhagic telangiectasia (HHT) or Osler-Weber-Rendu disease, is an autosomal dominant disease characterized by mucocutaneous telangiectasias, epistaxis and visceral arteriovenous malformations (AVM) most commonly found in the lungs, liver, and brain [1,2]. In Japan, the incidence of HHT is estimated to be 1:5000 to 1:8000 [3], which is similar to reports from other countries [4–6]. HHT is clinically diagnosed based on the Curaçao criteria: 1) an affected first-degree family member, 2) recurrent epistaxis, 3) multiple telangiectasia along the mucocutaneous surface, and 4) arteriovenous malformations in major organs. The diagnosis is considered “confirmed” in an individual with at least three, and “suspected” with two of the above features [4]. These features progress with age and pulmonary arteriovenous malformation (PAVM) development is thought to be complete by the end of puberty [7,8].

Recently, genetic research has demonstrated that heterozygous mutations including *ENG*, *AVCRL1*, and rarely *SMAD4* are causative

genes of HHT. There are at least two other unidentified genes that can cause HHT [7,8]. The majority of HHT patients have mutations in *ENG*, encoding endoglin, or *ACVRL1*, encoding activin receptor like kinase. These genes are associated with the transforming growth factor (TGF)- β superfamily signaling pathway, which is important for maintaining vascular integrity [5–7]. HHT with *ENG* mutation is characterized by a high incidence of PAVMs and cerebral AVMs [9], whereas HHT with *ACVRL1* is associated with hepatic AVMs. It is thought that clinical profiles have some correlation with the genotype.

Here we report four patients with familial HHT with PAVM associated with a new *ENG* mutation.

2. Case reports

In August 2015, a 48-year-old woman with HHT visited Sapporo Medical University Hospital for clinical genetic counseling with her four children, a son (Case 1) and three daughters (Case 2, 3, and 4). She had undergone left lower lobe lobectomy of a PAVM in the past. After

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Table 1
Summary of four cases.

Case	Age	Sex	Curaçao criteria	Symptom	PAVM			Treatment	Past history/Complications	
					Type	n	Location			Size (mm)
1	22	M	confirmed	Dyspnea Epistaxis	Simple	1	Right S10	60	Surgery	BA, ADHD, LD, Dysgraphia
2	27	F	confirmed	Dyspnea Epistaxis	Diffuse	3	Right S4 Right S6 Left S4	5 7 5	Surgery	BA, Irritable bowel syndrome
3	20	F	suggested	Dyspnea	Diffuse	14	Both lungs	2–18	TE	BA, ITP, ASD, Mental retardation, Depression, Mental intellectual disorder
4	12	F	confirmed	Epistaxis	Diffuse	3	Both lungs	2.8–6	none	Adjustment disorder, Mental disorder

BA: bronchial asthma, ADHD: attention deficit hyperactivity disorder, LD: learning disabilities, ITP: idiopathic thrombocytopenic purpura, ASD: autistic spectrum disorder, TE: transcatheter embolization.



Fig. 3. Case 1. Three-dimensional computed tomography (3D-CT) angiography reveals a large pulmonary arteriovenous malformation (PAVM, arrow) located in the subpleural area of the right segment (S) 10. The size is 60 mm in diameter and feeding pulmonary artery is 10mm in diameter.

episodes of epistaxis.

3. Discussion

We report a family with HHT with PAVMs associated with a new *ENG* mutation. Although more than a hundred mutations in the *ENG* gene have been reported related to HHT [6], there is no report of HHT caused by the *ENG* c. 1517T > C (p. Leu506Pro). To our knowledge, this is the first reported case. We would have been able to confirm whether this mutation was causative if there was an unaffected sibling without the mutation. Further research is needed in order to determine precisely if this mutation can cause HHT. In addition, their clinical profiles showed all of our patients had PAVM and three had psychiatric or developmental disorders. Recently, it was reported that pediatric patients with HHT had a relatively high prevalence of malformations of cortical development [10]. There may be a relationship between the

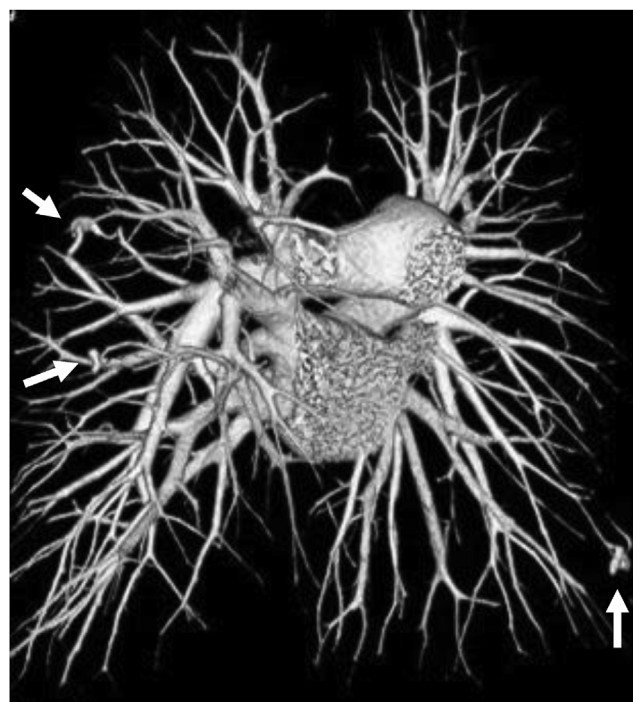


Fig. 4. Case 2. 3D-CT angiography reveals PAVMs (arrows) located in right middle and lower lobe, and left upper lobe. They are located in right S4, S6, and left S4. The sizes of them are 5 mm, 7 mm and 5 mm in diameter, respectively. They are simple type and supplied by a feeding artery of 3–4 mm in diameter.

observed psychiatric and developmental disorders and this new *ENG* mutation.

Causative genes of HHT including *ENG*, *ACVRL1* and *SMAD4* are involved in the TGF- β pathway [11,12]. The TGF- β superfamily signaling pathway has a crucial role in regulating proliferation, differentiation, and migration in angiogenesis. HHT gene mutations may negatively influence some forms of angiogenesis, with specific effects on the stability of newly formed vascular sprouts [7]. Endoglin, a membrane protein expressed in the endothelial cells, works as an endothelial specific co-receptor for multiple receptor complexes of the TGF- β superfamily. It has been hypothesized that the mutated proteins may be misfolded and unstable, and fail to reach the cell surface, rendering them unstable to form heterodimers with normal endoglin [6]. *ENG* mutations causing endoglin haploinsufficiency are thought to impair recruitment of mural cells to vessels [7].

Currently, transcatheter embolization with vascular coils and occluders (Amplatzer™) are the standard treatment for PAVMs [12]. Treatment of PAVM with transcatheter embolization of the feeding vessels significantly decreases right-to-left shunting, leading to



Fig. 5. Case 3. 3D-CT angiography reveals multiple PAVMs located in bilateral peripheral lung. The largest PAVM (arrow) with complex type is located in right S10.

improved oxygenation. In our cases, we performed this procedure for Case 1 and Case 2, and embolotherapy for Case 3. In Case 1, because of a large PAVM with thick feeding artery and draining vein which had presumably high flow of blood within the PAVM, endovascular treatment of the large PAVM was thought to have a higher risk of complications such as device embolism or thrombus passing through the PAVM. Akiyama et al. [13] reported a surgically treated case of PAVM who had turbulent thrombi in one of the PAVMs, which was capable of inducing embolic strokes such as symptomatic paradoxical thrombosis. Therefore, segmentectomy was thought to be preferable to transcatheter embolization in Case 1 for preventing severe complications with embolotherapy [13,14]. In Case 2, surgery was performed because embolotherapy was not feasible due to allergy to contrast medium. On the other hand, there is no agreement regarding the treatment of PAVM in pediatric patients [15]. A significant concern in treating growing lungs is that treated PAVMs may be at increased risk for reperfusion via pulmonary collaterals. Therefore, we determined to follow up Case 4 who was a 12-year-old child with asymptomatic multiple small PAVM. On the other hand, Case 1 showed recurrence and progression of PAVMs after treatment. The manifestations of progressing HHT and PAVM may take various forms with increasing age. Long-term follow-up is necessary for HHT patients even after AVM treatment.

Genetic testing is not always essential to diagnose of HHT, however, this knowledge might be useful for a patient with suggestive clinical features of HHT. In addition, when the affected parent has the mutation, the child without the mutation need not to undergo further screening test for AVM. In our cases, Case 3 was thought to be a pre-symptomatic mutation carrier. In this report, we presented 4 cases of HHT in a single family, in which the clinical diagnosis was confirmed using genetic testing, providing an opportunity for early detection of complications and management of the HHT.

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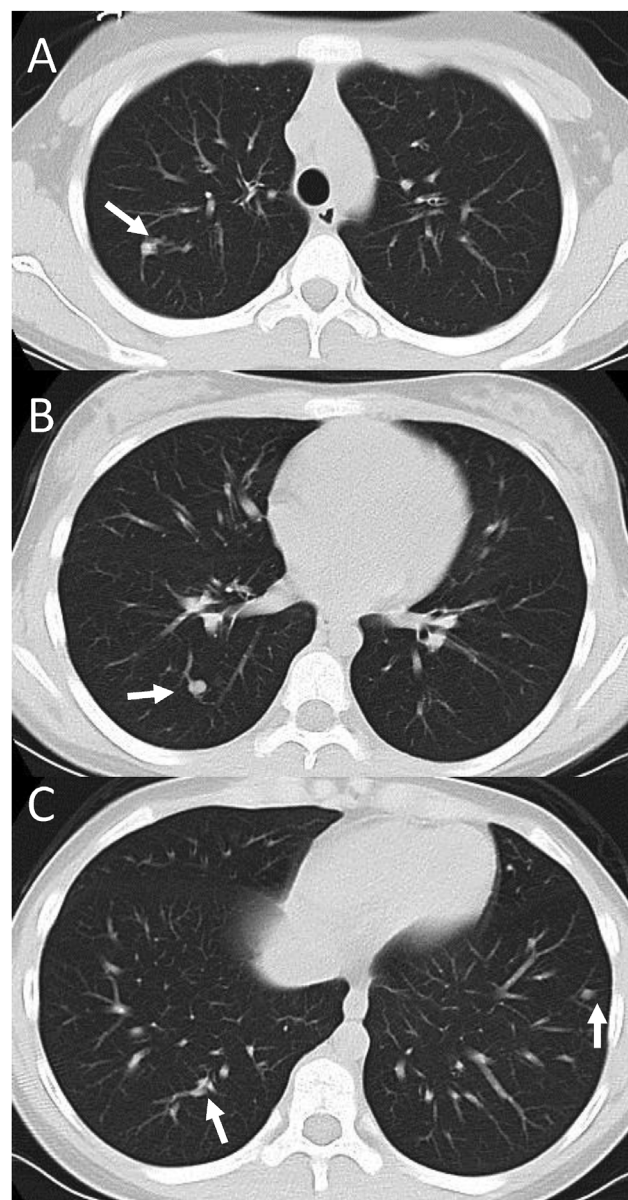


Fig. 6. Case 4. Chest CT shows three PAVMs (arrows) in right upper lobe (A), lower lobe (B) and left lower lobe.

Conflicts of interest

The authors state that they have no conflict of interest.

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