

[ CASE REPORT ]

## A Japanese Patient with Hereditary Myopathy with Early Respiratory Failure Due to the p.P31732L Mutation of Titin

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### Abstract:

Hereditary myopathy with early respiratory failure (HMERF) is caused by titin A-band mutations in exon 344 and is considered quite rare. Respiratory insufficiency can be the sole symptom in the disease course. We herein report the first Japanese HMERF patient with a p.P31732L mutation in titin. The patient manifested respiratory failure and mild weakness of the neck flexor muscle at 69 years old and showed fatty replacement of the bilateral semitendinosus muscles on muscle imaging. Our case indicates that HMERF with a heterozygous p.P31732L mutation should be included in the differential diagnosis of muscular diseases presenting with early respiratory failure.

**Key words:** hereditary myopathy with early respiratory failure (HMERF), myofibrillar myopathy, semitendinosus muscle, cytoplasmic body

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### Introduction

Respiratory muscle involvement in adults with muscular diseases usually occurs during the advanced stages of such disease and is uncommon at the disease onset. However, various adult-onset hereditary myopathies, including hereditary myopathy with early respiratory failure (HMERF), can be associated with respiratory muscle involvement while patients are still ambulatory (1-3).

Mutations in exon 344 encoding the 119th fibronectin-3 (FN3) domain in the A-band region of the *titin* gene are associated with HMERF (3-5). The reported mutations mainly show an autosomal dominant inheritance pattern, and only the p.P31732L mutation has been recognized as a semi-recessive or semi-dominant variant (5), showing incomplete penetrance (6).

We herein report the clinical features and histopathological and muscle computed tomography (CT) findings of the first Japanese HMERF patient with a heterozygous p.P31732L mutation.

### Case Report

The proband was a 69-year-old man. None of his family members, including his parents, siblings and children, had signs of muscle disease (Fig. 1). At 67 years old, he started experiencing dyspnea while farming, which necessitated frequent rests. In the same year, he was found to have atrial fibrillation (AF) without cardiomyopathy. He underwent catheter ablation for AF, leading to the termination of the tachycardia and sinus rhythm.

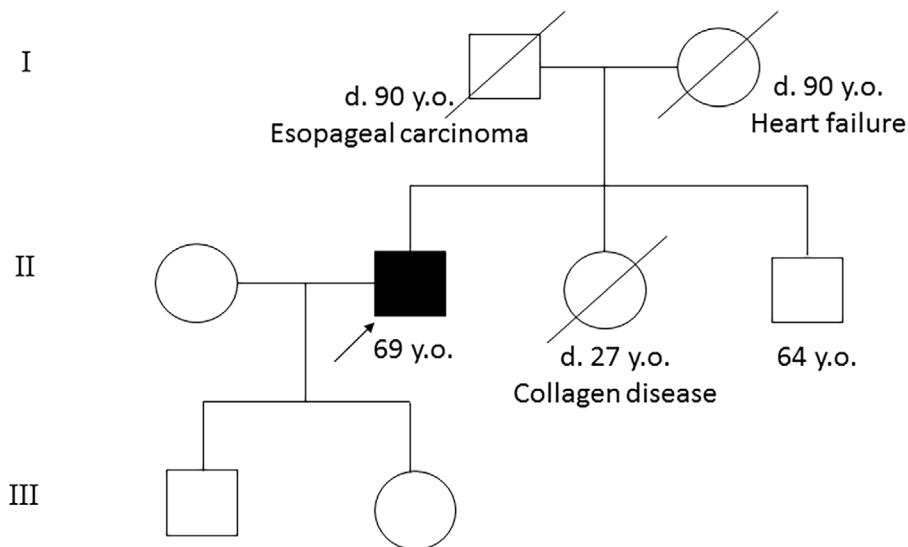
At 68 years old, he began to have daytime drowsiness. At 69 years old, he felt heart palpitations and fatigue and was diagnosed with heart failure with a low cardiac output due to atrial flutter (AFL). High-density oxygen was administered, and his consciousness level gradually decreased. Thereafter, he presented with respiratory failure with hypercapnic coma and required mechanical ventilation and tracheotomy.

After recovering from the heart failure and AFL, he was admitted to our department for a further investigation regarding a possible neurological cause of his restrictive pul-

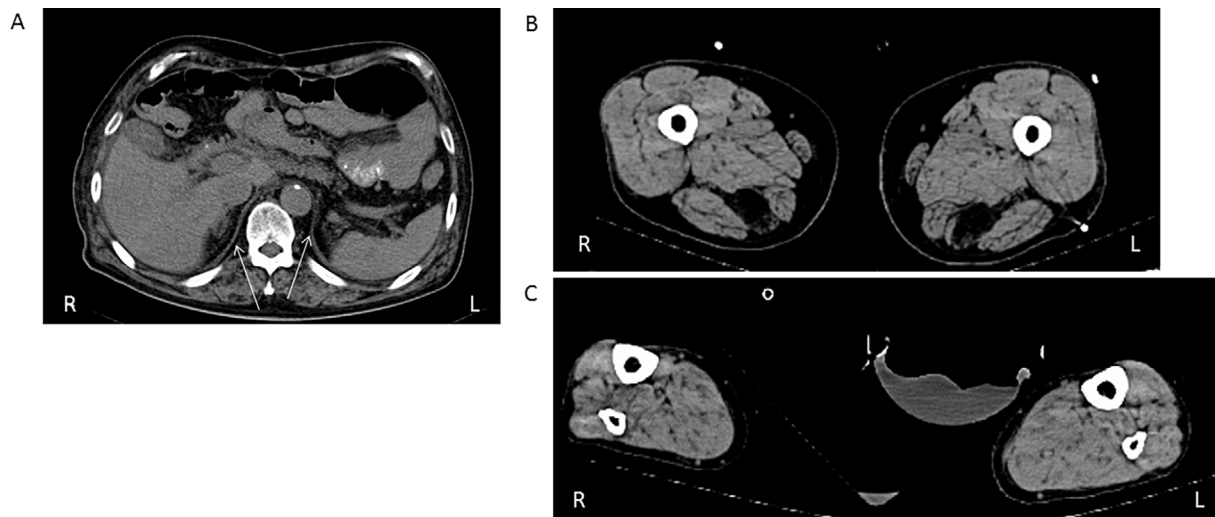
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**Figure 1.** The parents of the proband are not consanguineous. The squares and circles indicate males and females, respectively. The proband is indicated by an arrow and is the only person with this disease in this pedigree. Diagonal lines through symbols represent deceased persons with the cause of death indicated for each person.

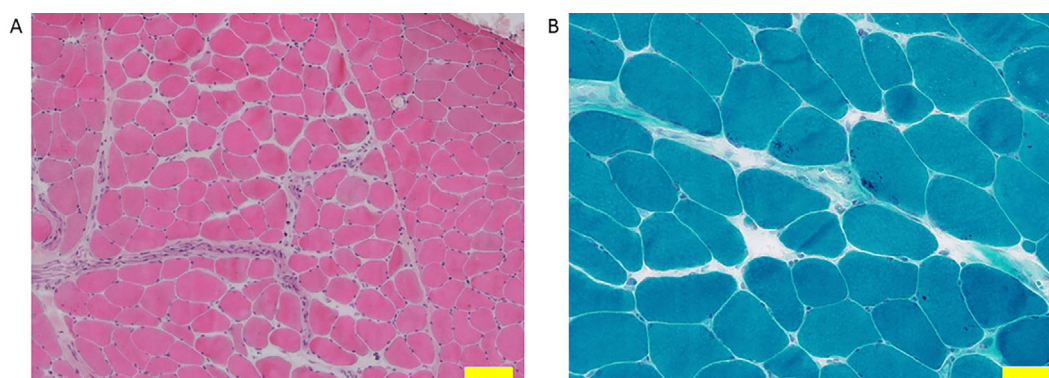


**Figure 2.** Muscle CT findings of the proband. A: On abdominal CT, atrophy of the diaphragm (arrow) can be seen. B: At the thigh level, fatty degenerative changes in the semitendinosus muscles are noted. C: At the lower leg level, anterior lower muscles, including peroneus longus, are preserved.

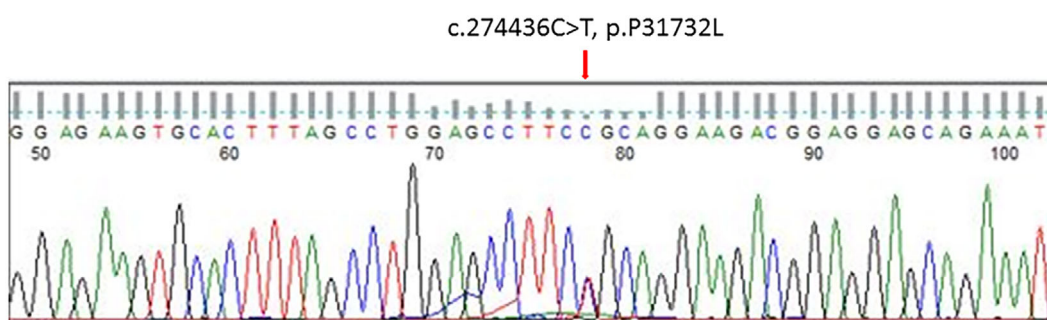
monary disease. Although mild weakness of the neck flexor muscles was noted, he had full muscle strength of the upper and lower limbs. His gait was normal before admission, as was his serum creatine kinase level. Chest expansion on respiration was poor, suggesting respiratory muscle involvement, and his forced vital capacity was 61%. An electromyography (EMG) examination of the right semitendinosus muscle showed myopathic abnormalities. Nerve conduction studies were normal. Muscle CT revealed atrophy of the diaphragm (Fig. 2A) and fatty replacement of the bilateral semitendinosus muscles (Fig. 2B). The tibialis posterior muscles were preserved (Fig. 2C). A muscle biopsy performed on the left biceps brachii muscle showed myopathic features, such as increased fiber size variation on hematox-

ylin and eosin staining (Fig. 3A) and cytoplasmic bodies (CBs) on modified Gomori trichrome staining (Fig. 3B).

He was diagnosed with myofibrillar myopathy (MFM) according to the clinical feature and the CBs on a muscle biopsy. Paired-end sequencing of exons of the genes to screen for MFM, including *ACTA1*, *BAG3*, *CFL2*, *CRYAB*, *DES*, *DNAJB6*, *EPG5*, *FHL1*, *FLNC*, *GNE*, *KBTBD13*, *KLHL40*, *LAMP2*, *LDB3*, *MATR3*, *MEGF10*, *MYH2*, *MYH7*, *MYOT*, *NEB*, *ORAI1*, *PABPN1*, *PLEC*, *RBCK1*, *SEPN1*, *SIL1*, *STIM1*, *TCAP*, *TIA1*, *TNNT1*, *TPM2*, *TPM3*, *TRIM32*, *TTN*, *VCP* and *VMA21*, was performed using a next-generation Ion PGM™ sequencer (Thermo Fisher Scientific, Carlsbad, USA) (7). The variants were filtered by allele frequency using public databases, including 1,000 genomes, the Exome



**Figure 3.** Histopathological features. A: Hematoxylin and Eosin staining shows increased fiber size variation, indicating a myopathic feature (scale bar: 100 microns). B: A modified Gomori trichrome shows cytoplasmic bodies in the subsarcolemmal position (scale bar: 50 microns).



**Figure 4.** Sanger sequencing confirmed the heterozygous missense mutation [c.274436C>T, p.P31732L] of the proband.

Aggregation Consortium and ClinVar. To predict the disease-causing potential for variants, we used Mutation Taster and PolyPhen2, and to evaluate the pathogenicity of the causative variants, we used the American College of Medical Genetics (ACMG) 2015 guideline (8).

Finally, we identified a heterozygous missense variant in *TTN* on chromosome 2:178546041 in reference to the Genome Reference Consortium Human Genome Build 38 (GRCh38). This variant is listed in gnomAD at an allele frequency of  $1.21 \times 10^{-5}$ . According to 8.3KJPN, this variant was not identified in 8,300 Japanese individuals. In addition, this variant was classified as a ‘pathogenic variant’ by the ACMG guideline. This variant [g.274436C>T, (c.95195C>T; p.P31732L), NM\_001267550.2] was confirmed by Sanger sequencing (Fig. 4). Target re-sequencing did not reveal any variant in the kinase domain of *TTN*, including p.R34091W, which has been reported to be linked to p.P31732L in European patients (9). The patient continued invasive nocturnal ventilation support thereafter and was transferred to another hospital for rehabilitation of his respiratory muscles.

### Ethics

All experiments in this study were approved by the Ethics Committees of Yamaguchi University Hospital and the National Center of Neurology and Psychiatry. Written informed consent was obtained from the proband.

## Discussion

Respiratory muscle weakness occurs in a broad variety of neuromuscular diseases during the late stage of the disease. However, this condition can also occur while patients are still ambulatory. Amyotrophic lateral sclerosis (10) and myasthenia gravis (11) are important diseases to consider, but even after these have been excluded, the differential diagnosis of adult-onset hereditary myopathies remains challenging. Naddaf et al. reported that MFM was the second-most common condition after adult-onset Pompe disease among hereditary myopathies as a cause of early respiratory failure (12). Besides HMERF, there have been case reports of patients presenting with respiratory failure in MFM due to mutations in *DES* (13), *CRYAB* (14) and *BAG3* (15). We should therefore consider MFM as an important differential diagnosis of diseases manifesting as early respiratory insufficiency.

The importance of neuromuscular imaging in recognizing the degree and pattern of muscle involvement is increasing, helping to limit the range of the differential diagnosis and facilitating making the actual diagnosis. The predominant involvement of the semitendinosus on imaging, or even its isolated involvement, is considered specific to a group of MFMs, including HMERF and those caused by mutations in

**Table 1. Japanese Patients with HMERF.**

Patient	Age/ Sex	Mutation (protein level)	Initial manifestation (age)	Respiratory disturbance (age)	Selective muscle involvement on imaging		Foot drop	References
					ST	ACLs		
A-1	49/M	p.C31712R	Tripping (46)	Yes (48)	NA	NA	+	19
A-2 (Son of A-1)	26/M	p.C31712R	Tripping (20)	Yes (*)	+	+	+	19
B-1	45/M	p.C31712R	Foot drop (31)	Yes (44)	+	+	+	19
B-2 (Younger brother of B-1)	37/M	p.C31712R	Foot drop (27)	Yes (34)	+	+	+	19
C	34/F	p.C31712R	Fatigability (26)	Yes (26)	+	+	+	19
D	38/M	p.C31712R	Difficulty in lifting theigh (20)	Yes (26)	+	-	+	19
E	40/M	p.C31712R	Fatigability, respiratory failure (31)	Yes (31)	+	+	+	19
F	52/M	p.C31712R	Foot drop (47)	Yes (50)	+	+	+	19
G-1	67/M	p.C31712R	Tripping (57)	Yes (*)	+	+	+	19, 22
G-2 (son of G-1)	38/M	p.C31712R	Tripping (20)	Yes (29)	+	+	+	19, 22
G-3 (daughter of G-1)	36/F	p.C31712R	Gait disturbance (31)	Yes (34)	+	+	+	24
H	68/F	p.C31712R	Difficulty in standing on right toe (68)	Yes (68)	+	+	-	19
I	43/M	p.C31712R	Fatigability (39)	Yes (39)	+	+	-	19
J	42/F	p.C31712Y	Difficulty in lifting theigh (36)	Yes (*)	+	+	-	19
K	38/M	p.G31791D	Gait disturbance (31)	Yes (*)	+	+	+	19
L	44/F	p.G31791R	Fatigability (26)	Yes (40)	+	+	-	19
M	40/M	p.G31791V	Gait disturbance (24)	Yes (27)	**	+	+	19
N	46/M	p.R31783_ V31785del	Foot drop, difficulty in opening a bottle (41)	Yes (41)	+	+	+	19
O	69/M	p.P31732L	Respiratory failure (67)	Yes (67)	+	-	-	Present patient

NA: not available, ST: semitendinosus, ACLs: anterior compartment of lower legs

\*Asymptomatic but found by laboratory tests

\*\* Diffuse muscle involvement

*DES* and *CRYAB* (4, 16, 17). However, we ruled out mutations in the exons of both of *DES* and *CRYAB* by the panel analysis.

HMERF is distributed worldwide, including in Europe, North and South America and Asia (18, 19). Several dominant mutations in the *TTN* A-band have been identified in this disease, including the c.95134T>C, p.C31712R, which is the most frequent HMERF mutation in Europe (18) and Japan (Table 1). Among the mutation patterns, there is still some debate concerning the nature of the p.P31732L mutation (Table 2). Among 14 individuals who were p.P31732L heterozygotes or homozygotes, 3 (B-I-1, B-I-2 and D-II-2) showed no signs of muscle disease, two (A-I-1 and A-I-2) showed a mild subclinical phenotype with typical muscle magnetic resonance imaging (MRI) findings, 3 (C-II-1, E-II-1 and E-II-2) showed only respiratory failure, 2 (C-II-2 and G-II-1) - including our patient - showed only neck flexor weakness with respiratory failure, 2 (D-II-1 and F-II-1) showed neck flexor and limb weakness as well as respiratory failure, and 2 homozygotes (A-II-1 and B-II-1) had more severe disease with an earlier disease onset (Table 2).

Although whether or not selective muscle involvement on imaging was seen in unaffected p.P31732L carriers (B-I-1, B-I-2, and E-II-2) was not mentioned, it remains possible that the p.P31732L mutation is a pathogenic variant with variable penetrance.

In contrast, Lange et al. suggested that the p.P31732L mutation is recessive and can become penetrant when the *in cis* *TTN* kinase variant p.R32450W works in synergy on the protein turnover pathways (9). However, the p.P31732L mutation was identified as the cause of HMERF in other families, without any concomitant *TTN* kinase variant (5, 6, 20). In addition, Yue et al. reported a 23-year-old Chinese HMERF patient with a heterozygous p.P31732L mutation without an accompanying kinase domain mutation (21). Furthermore, our patient also had no mutation in the kinase domain of *TTN*. These findings indicate that the heterozygous p.P31732L mutation itself is a cause of HMERF, although the phenotype of this mutation is definitely variable.

Muscle weakness in HMERF together with early respiratory failure usually involves the proximal, distal and trunk muscles as the disease progress, and the onset is between

**Table 2. Clinical Features of Individuals Having p.P31732L Mutation.**

Patient	Age/ sex	Initial manifestation (age)	Muscle weakness	Respiratory disturbance	Selective muscle involvement on imaging		Nationality	References
					ST	ACLLs		
A-I-1 (father of A-II-1)	59/M	-	-	-	+	-	Italian	5
A-I-2 (mother of A-II-1)	56/F	-	-	-	+	-	Italian	5
A-II-1 (Ho)	32/M	Nocturnal hypoventilation, exertional dyspnea (30)	Neck flexor, proximal LL, ankle dorsiflexion	+	*	*	Italian	5
B-I-1 (father of B-II-1)			-	-	NA	NA	French**	5
B-I-2 (mother of B-II-1)			-	-	NA	NA	French**	5
B-II-1 (Ho)	36/M	Effort breathlessness (27)	proximal UL, proximal LL, finger extensors, distal LL, wheelchair bound	+	*	*	French**	5
C-II-1	56/M	Respiratory failure (42)	-	+	NA	-	French**	5
C-II-2	58/F	Hypoventilation (53)	Neck flexor	+	NA	NA	French**	5
D-II-1	57/M	Difficulty in lifting LL (30s)	Neck flexor, proximal UL proximal LL	+	NA	NA	British	6
D-II-2 (younger brother of D-II-1)		-	-	-	NA	NA	British	6
E-II-1	56/M	Respiratory failure (46)	-	+	NA	NA	Portuguese	20
E-II-2 (sister of E-II-1)		NA	-	+	NA	NA	Portuguese	20
F-II-1	23/M	Difficulty in bending the neck (23)	Neck flexor Knee flexor	+	+	+	Chinese	21
G-II-1	69/M	Respiratory failure (67)	Neck flexor	+	+	-	Japanese	Present patient

NA, not available. sT, Semitendinosus. Ho: homozygote

Only A-II-1 and B-II-1 are homozygotes, and others are heterozygous state with p.P31732L.

\* Diffuse muscle involvement

\*\* with Portuguese ancestry

the second and the fifth decades of life (2-4). Almost all Japanese patients with HMERF also show an onset before the fifth decade of life and have limb weakness (Table 1). However, the onset of disease in the present patient occurred at 67 years old, which is very late among HMERF individuals, and only weakness of the neck flexor muscles manifested, aside from respiratory muscle involvement. In their first description of HMERF, Edström et al. reported that all patients were characterized by proximal muscle weakness of the upper and lower extremities, with early affection of the neck flexors and respiratory muscles, especially the diaphragm (2). A late-onset HMERF patient like our own might therefore only manifest respiratory failure and mild neck flexor weakness as a clinical presentation. In addition, among seven clinically affected p.P31732L heterozygotes, five only showed respiratory failure with or without neck

flexor weakness (Table 2). Heterozygous patients with this peculiar mutation of TTN might therefore manifest a milder phenotype than those with other dominant mutations with the FN3 domain of the TTN A-band.

In muscle specimens of HMERF, CBs are often located in the subsarcolemmal region, resulting in their being referred to as 'necklace CBs' (5, 19). Uruha et al. reported that necklace CBs were found in 14 of 17 patients with genetically-confirmed HMERF (19). Based on the result, the sensitivity of the necklace CBs in HMERF was calculated to be 82%. The CBs in our patient did not show a necklace-like alignment, possibly due to the older onset and milder presentation or to the fact that the biopsy was not properly targeted. In either case, these findings suggest that HMERF should be considered in patients associated with both early respiratory failure and non-specific CBs, as in our patient.

In conclusion, HMERF with a p.P31732L mutation should be included as a differential diagnosis for ambulatory patients with respiratory insufficiency and neck flexor weakness who harbor non-specific CBs on muscle biopsy specimens.

**The authors state that they have no Conflict of Interest (COI).**

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