

ORIGINAL ARTICLE

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Mitochondrial DNA mutations in Japanese patients with optic neuropathy unassociated with a mutation at nucleotide position 11,778

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Abstract We examined for mitochondrial DNA (mtDNA) mutations at nucleotide positions (nt) 3460, 14,484, 9438, 9804, and 15,257 in ten Japanese patients with idiopathic optic neuropathy unassociated with a mutation at nt11,778. The mtDNAs were amplified by polymerase chain reaction (PCR), the products were digested with restriction enzymes, and the sizes of the fragments were analyzed on 8% polyacrylamide gel. Of the ten patients, one had an mtDNA mutation at nt3460 and another patient had a mutation at nt14,484. We suggest that mtDNA mutations in Japanese patients with optic neuropathy unassociated with a mutation at nt11,778 should be further investigated.

Key words Leber hereditary optic neuropathy · nt3460 mutation · nt14,484 mutation · Optic neuropathy · Japanese patients

Introduction

Leber hereditary optic neuropathy (LHON) is a maternally inherited disease characterized by acute visual impairment in both eyes. The disease is associated with 17 missense mutations in mitochondrial DNA (mtDNA) (McKusick 1994). Of the 17 mutations, 3 are thought to be primary, and the presence of these three greatly increases the possibility of blindness (Brown et al. 1997). Mutations in mtDNA have been commonly reported at nucleotide positions (nt) 11,778, 3460, 14,484, 9438, 9804, and 15,257 (Wallace et al.

1988; Fujiki et al. 1991; Huoponen et al. 1991; Majander et al. 1991; Johns et al. 1992a and 1992b; Johns and Neufeld 1993; Mashima et al. 1993; Newman 1993; Nakamura et al. 1994; Oostra et al. 1994; Hotta et al. 1995; Riordan-Eva et al. 1995; Fujimaki et al. 1996; Ghosh et al. 1996; Joo et al. 1996; Nikoskelainen et al. 1996). A high frequency of mtDNA mutations at nt11,778 has been reported in Japanese patients with the disease (Fujiki et al. 1991; Mashima et al. 1993; Hotta et al. 1995). In an earlier study, we examined for the nt11,778 mutation in 39 Japanese patients with idiopathic optic neuropathy, and reported clinical features in 29 LHON patients with the nt11,778 mutation (Fujimaki et al. 1996). In the present study, we examined for mtDNA mutations at nt3460, 14,484, 9438, 9804, and 15,257 in the 10 Japanese patients with idiopathic optic neuropathy unassociated with the nt11,778 mutation in that earlier study.

Subjects and methods

In an earlier study (Fujimaki et al. 1996), we examined for the nt11,778 mutation in mtDNA in 39 individuals with idiopathic optic neuropathy who were patients at the Eye Clinic of Juntendo University Hospital from 1989 to 1995. Twenty-nine of the 39 patients had the nt11,778 mutation. In the present study, we examined the 10 remaining patients whose idiopathic optic neuropathy was unassociated with the nt11,778 mutation. Informed consent for the present study was obtained from each patient.

mtDNA was extracted from cells of peripheral blood samples. We used five pairs of primers to determine nt3460, 14,484, 9438, 9804, and 15,257 mutations (Table 1). The DNA fragments were amplified by PCR, using a DNA Thermal Cycler (Perkin-Elmer Cetus, Branchburg, NJ, USA). The purified PCR products were digested with restriction enzymes (New England Bio Labs, Beverly, MA, USA and Boehringer Mannheim, Mannheim, Germany) (Table 1), and the sizes of the fragments were determined on 8% polyacrylamide gel containing ethidium bromide.

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Table 1 Polymerase chain reaction primers for mtDNA mutations

Nucleotide position	Primers	Restriction enzyme
11,778 (G to A)	Sense: ATACTCTTCAATCAGCCACATAGC Antisense: GAGAACGTGGTTACTAGCACAGAG	<i>Sfa</i> N I and <i>Mae</i> III
3460	Sense: CACACCCACCCAAGAACAGGG Antisense: GAGATTGTTTGGGCTACTGC	<i>Hga</i> I
14,484	Sense: AGTATATCCAAAGACAACGA ^a Antisense: GATGTATGCTTTGTTTCTGT	<i>Sau</i> 3A I
9438	Sense: ATCCAAGCCTACGTTTTC Antisense: GATGAAGCAGATAGTGAG	<i>Stu</i> I
9804 (G to A)	Sense: ATCCAAGCCTACGTTTTC Antisense: GATGAAGCAGATAGTGAG	<i>Mae</i> III
15,257	Sense: CTCCTGCTTGC AACTAT Antisense: CCGAGGGCGTCTTTGAT	<i>Acc</i> I

mt DNA, Mitochondrial DNA

^a The underline indicates a mismatched nucleotide

Table 2 Clinical features and mtDNA mutations in patients with optic neuropathy

Patient no.	Age at onset (years)	Sex	Family history	Findings at initial visit				mtDNA mutations					
				Visual acuity		Optic disc		11,778	3460	14,484	9438	9804	15,257
				OD	OS	OD	OS						
1	19	M	+	0.08	0.02	P	P	–	–	–	–	–	–
2	17	F	–	0.4	0.7	P	P	–	+	–	–	–	–
3	20	M	–	0.09	0.06	P	P	–	–	+	–	–	–
4	39	M	–	0.04	1.0	S	N	–	–	–	–	–	–
5	10	M	–	0.3	0.2	P	P	–	–	–	–	–	–
6	37	M	–	0.15	0.1	P	P	–	–	–	–	–	–
7	12	M	–	CF	1.0	S	N	–	–	–	–	–	–
8	17	M	–	0.08	0.08	P	P	–	–	–	–	–	–
9	13	M	–	0.5	0.5	P	P	–	–	–	–	–	–
10	19	M	–	0.1	HM	P	P	–	–	–	–	–	–

CF, Counting fingers; HM, hand motion; P, pallor; S, swelling; N, normal; +, positive; –, negative; OD, right eye; OS, left eye

Results

The clinical features and mtDNA mutations in the ten patients with optic neuropathy are shown in Table 2. Eight patients had bilateral optic disc pallor and two had unilateral optic disc swelling. Patient 1 had a family history of optic atrophy: his maternal uncle had bilateral optic atrophy. This patient, however, had no mutation at either nt11,778, 3460, 14,484, 9438, 9804, or 15,257. Patient 2 was diagnosed with bilateral optic atrophy at age 17 years. She had relatively good visual acuity in both eyes (0.4 OD and 0.7 OS). She had no family history of optic neuropathy, but she had an mtDNA mutation at nt3460 (Fig. 1). Patient 3 was diagnosed with bilateral optic atrophy at age 20 years. He had poor visual acuity in both eyes (0.09 OD and 0.06 OS). He had no family history of optic neuropathy, but he had a mutation at nt14,484 (Fig. 2). Patients 4–10 had no family history of optic neuropathy and no mutations at nt11,778, 3460, 14,484, 9438, 9804, or 15,257.

Discussion

In our previous study of 39 patients with LHON and with possible hereditary optic neuropathy, we found that 29

had an mtDNA nt11,778 mutation (Fujimaki et al. 1996). In the present study of the remaining 10 patients in whom the optic neuropathy was unassociated with the nt11,778 mutation, patient 2 had an mtDNA mutation at nt3460, and patient 3 had an mtDNA mutation at nt14,484.

Nakamura et al. (1994), Hiida et al. (1995), and Joo et al. (1996) reported an nt3460 mutation in Japanese patients with LHON and suggested that the prevalence of the mutation in Japanese patients may be lower than that in patients of other ethnic origins. However, it is likely that the prevalence of the nt3460 mutation in Japanese patients with LHON may be higher than previously thought, in the light of our finding of one patient with this mutation in the present screening of a small number of patients with idiopathic optic neuropathy. Patients with LHON and the nt3460 mutation are thought to have better visual function than those with the nt11,778 mutation (Johns et al. 1992b; Joo et al. 1996; Nakamura et al. 1994; Newman 1993; Nikoskelainen et al. 1996). Patient 2 in the present study also had relatively good visual acuity in both eyes. We were unable to examine for the mutation in her asymptomatic maternal relatives. Majander and co-workers (1991) reported that nicotinamide adenine dinucleotide, reduced (NADH): ubiquinone reductase had different electron transfer properties in the NADH dehydrogenase subunit 1

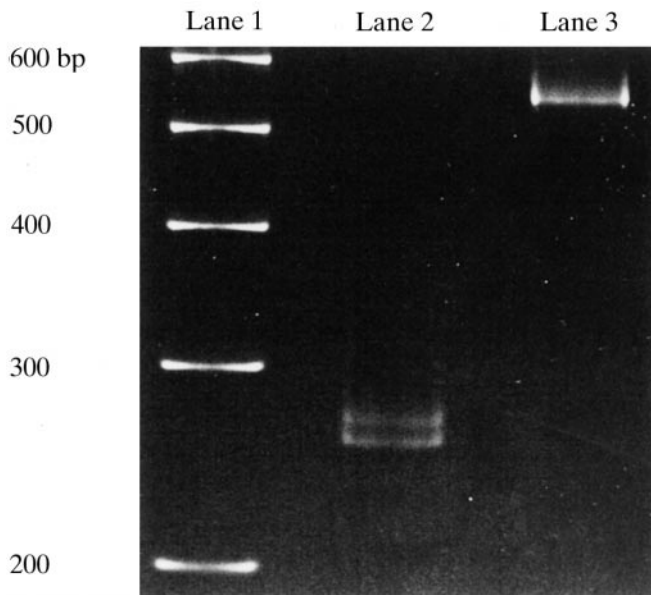


Fig. 1 Analysis of mutation at nucleotide position (nt)3460 of mitochondrial (mt)DNA. Shown is 8% polyacrylamide gel electrophoresis after *Hga* I digestion of polymerase chain reaction (PCR) products encompassing nt3460. Lanes 1, 2, and 3 represent size marker, normal control, and patient 2 sample, respectively. Numbers at left indicate base-pair lengths of corresponding bands. Note that lane 2 has two visible bands, of 254 and 268 base-pair lengths, by *Hga* I digestion. Lane 3 exhibits only a 522-base pair length band, due to loss of the enzyme's recognition site, demonstrating a mutated sequence in this region

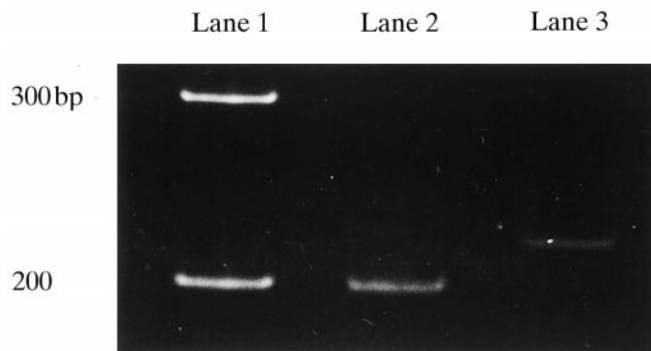


Fig. 2 Analysis of mtDNA mutation at nt14,484. Shown is 8% polyacrylamide gel electrophoresis after *Sau3A* I digestion of PCR products encompassing nt14,484. Lanes 1, 2, and 3 represent size marker, normal control, and patient 3 sample, respectively. Numbers at left indicate base-pair lengths of corresponding bands. Note that lane 2 has a band of 190-base pair length by *Sau3A* I digestion. Lane 3 exhibits a 208-base pair length band, due to loss of recognition site of *Sau3A* I, demonstrating a mutated sequence in this region

(ND1)/3460 and the ND4/11,778 mutations of LHON. These different biochemical properties may result in different clinical features.

Johns et al. (1992b), Oostra et al. (1994), and Riordan-Eva et al. (1995) reported an mtDNA mutation at nt14,484 in patients with LHON. Riordan-Eva et al. (1995) showed that good visual outcome was strongly correlated with age at onset in the patients with the nt14,484 mutation; in all

those whose onset was before age 20 years, final visual acuity was better than 6/24. Patient 3 in the present study developed optic neuropathy at age 20 years. Although Yamada et al. (1997) indicated that patients with the nt14,484 mutation had good visual outcome, our patient 3 had poor visual acuity. We were not able to examine for the mutation in his asymptomatic maternal relatives.

We suggest that mtDNA mutations in patients with idiopathic optic neuropathy unassociated with the nt11,778 mutation should be further investigated to enable precise diagnosis and appropriate genetic counseling.

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