

## Letter to the Editor

# **N98S mutation in NEFL gene is dominantly inherited with a phenotype of polyneuropathy and cerebellar atrophy**

**Keywords:**

Charcot–Marie–Tooth disease

NEFL

Dysmyelination

Axonal loss

Myelin

Nerve conduction study

Cerebellar atrophy

**1. Introduction**

Charcot–Marie–Tooth disease (CMT) is classified into either dysmyelinating or axonal type. The former exhibits slowed conduction velocities on nerve conduction studies (NCS) and abnormally developed myelin on nerve biopsies. In contrast, the latter shows normal or minimally slowed conduction velocities with axonal loss in its neuropathology. CMT can be transmitted in autosomal dominant, recessive or X-linked mode [3].

The *neurofilament light polypeptide* (NEFL) gene is located on 8p21. It encodes a neuronal protein that serves as a key element in the formation of neurofilaments and, therefore, affects axonal transport and nerve conduction velocity [4,6]. At least 20 missense mutations and a few truncation or frame-shift mutations in NEFL have been reported to cause either axonal CMT2E or dysmyelinating CMT1F (Fig. 1, upper panel). Patients may develop sensory loss and muscle atrophy/weakness in limbs.

Of these missense mutations, N98S mutation was only briefly described in *de novo* subjects. Thus, it is unknown if the mutation is heritable. Here, we report a CMT family with a dominantly inherited N98S mutation.

**2. Case report**

The proband (a 40-year-old woman) was an adopted child (Fig. 1B). She did not start walking until the age of 3 years and even then had a “drunken” gait and frequent falls. By the age of 8 years, she developed foot deformities and underwent bilateral Achilles tendon release. Throughout adolescence and early adulthood, she used ankle-foot orthoses to walk and experienced a continued progression of unsteady gait and falls. She acquired a scooter 4–5 years ago. Since 2 years ago, she has noted the inability to transfer herself out of the scooter.

Her neurological examination showed normal cognitive functions. She had a right eye esotropia and hearing loss. She had flat feet with hammer toes. Her muscle strength was 0–2 on MRC scale for distal muscles and 3–4 for proximal muscles. There was no truncal ataxia. Severe

limb weakness made the examination of coordination difficult. Her sensory functions were diminished to all modalities in her limbs. Tendon reflexes were absent. Her CMT Neuropathy score (CMTNS) was 31. The CMTNS is a validated measurement of impairment in patients with CMT [5] with a maximum of 36. It is classified as: the presence of mild (1–10), intermediate (11–20) or severe disability ( $\geq 21$ ). In addition, she had a brain MRI that revealed severe cerebellar and spinal cord atrophy (Fig. 1C).

The proband's son (8 years old) is her only biological child. He did not walk until 19 months. Frequent falls were noted throughout childhood and he struggled with manipulating buttons and shoelaces. He was also found to have mild, bilateral sensorineural hearing loss. Academic performance was reportedly normal. He had bilateral Achilles tendon release surgery at 7 years of age.

Neurological examination showed normal cognitive functions. He had high-arched feet. His muscle strength was 4/4 in bilateral ankle dorsal flexors but 5/5 in the remaining muscles. Sensation to all modalities was reduced in his feet. Tendon reflexes were absent.

NCS in the proband showed that all sensory nerves were non-responsive. CMAP was not recordable in distal muscles but was recordable in forearm muscles. The amplitude (0.5 mV) and conduction velocity (31 m/s) were reduced in the median nerve. CMAP was absent for the ulnar nerve. For the son, the sensory nerves were also non-responsive. The distal latency, amplitude and conduction velocity in the son's median, ulnar and peroneal nerves were 6.3 ms, 3.1 mV, 25 m/s; 4.0 ms, 6.0 mV, 31 m/s and 5.9 ms, 3.0 mV, 32 m/s, respectively.

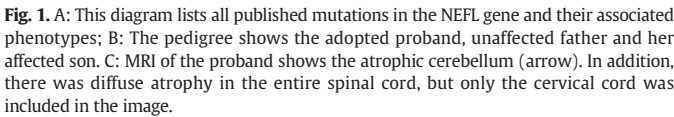
Their DNA tests were done by Medical Neurogenetics, Inc. in Atlanta. It demonstrated a N98S missense mutation of NEFL in the proband and the son. This study was approved by the Vanderbilt Institutional Review Board. The consent was obtained from the proband.

**3. Discussion and conclusion**

N98S mutation has been reported in four patients. All of them were *de novo* mutations with no detailed phenotypic description. It has been speculated that this mutation is embryonically lethal. The family in this report demonstrates that the N98S mutation is still heritable via an autosomal dominant mode.

This notion is further supported by another family that was reported shortly after our manuscript submission. A 6-year-old boy inherited a heterozygous N98S mutation from his affected mother [2]. However, in contrast to our study, the boy reported by Berciano et al. was mildly affected and the atrophy in the proband only affected cerebellum but not spinal cord. The two families together do not show obvious anticipation. In addition, sequencing of spinocerebellar ataxia (SCA) genes showed no mutation in the Berciano-reported family.

NEFL encodes the neurofilament light polypeptide with 543 amino acids. It provides structural support for the axon and regulates the axon caliber and transport [6]. The N98S mutation promotes protein aggregates in the cytoplasm. These aggregates appear to prevent NEFL from transporting to axons [1,4,8]. Indeed, decrease of neurofilament



A knockin mouse model of N98S has recently been produced [1]. In addition to the expected pathology, such as protein aggregates, in the peripheral nerves and spinal neurons, cerebellar neurons were predominantly affected in the central nervous system. Thus, the concordant findings between human and mouse support the utility of the knockin mouse.

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