





Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico

Sistema Socio Sanitario



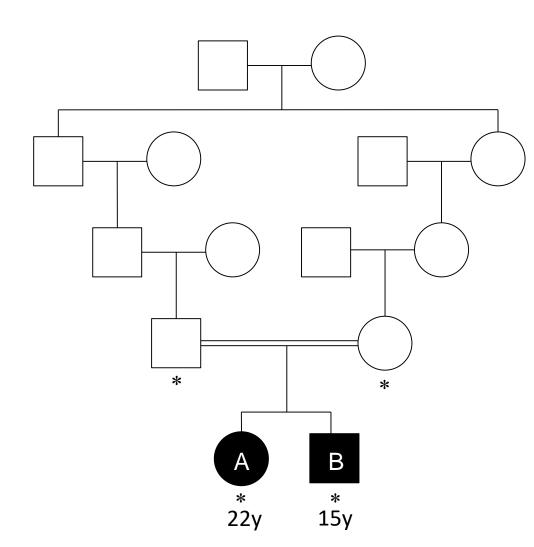
Neurofascin (*NFASC*) is a novel gene causing hereditary ataxia

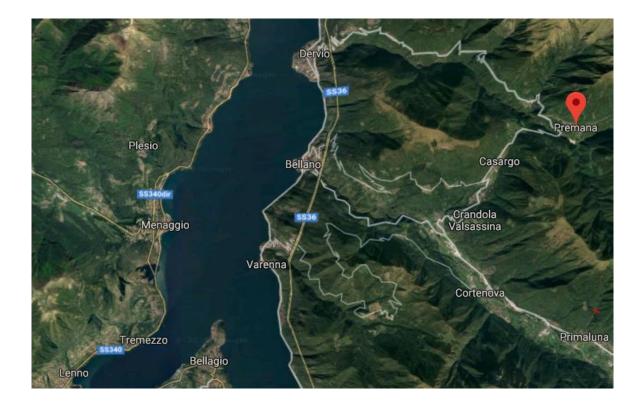
A family-based approach to find the genetic cause of a new neurological disease



Edoardo Monfrini Specializzando in Neurologia

Early-onset ataxia in an Italian consanguineous family





The family originated from an isolated valley of Lombardia region.

The parents are related (cousins)

Clinical features of the patients

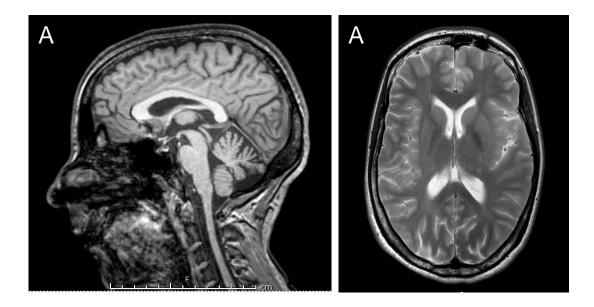
Early psychomotor delay was present. They started to walk at 3 years of age with an unsteady **wide-based ataxic** gait.

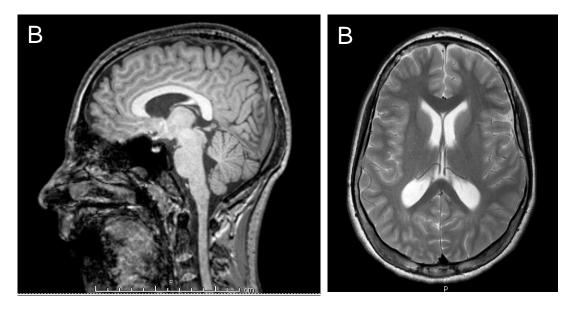
Prominent cerebellar signs: cerebellar dysarthria, intention tremor, dysmetria, pursuit saccadization, and dysdiadochokinesia were observed.

From the age of 11 years, subject A developed progressive spastic hypertonia of the limbs with hyperreflexia, lower limbs intra-rotation, ankle clonus, and bilateral Babinski sign.

Nerve conduction studies showed a length-dependent demyelinating sensory-motor peripheral neuropathy.

→ Known etiological genes for hereditary ataxias were excluded.

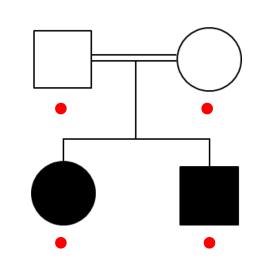




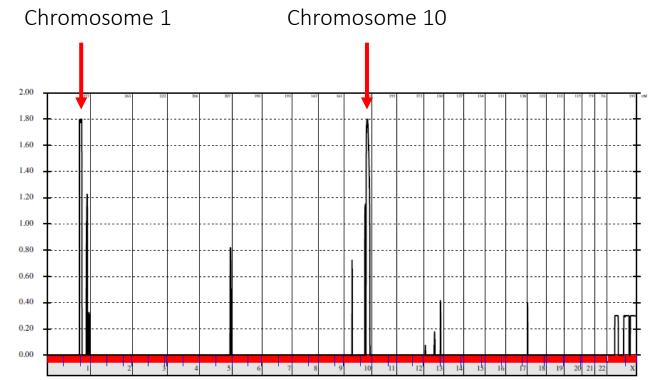
Linkage analysis – Homozygosity mapping

A homozygous mutation was suspected to be the cause of disease in this family



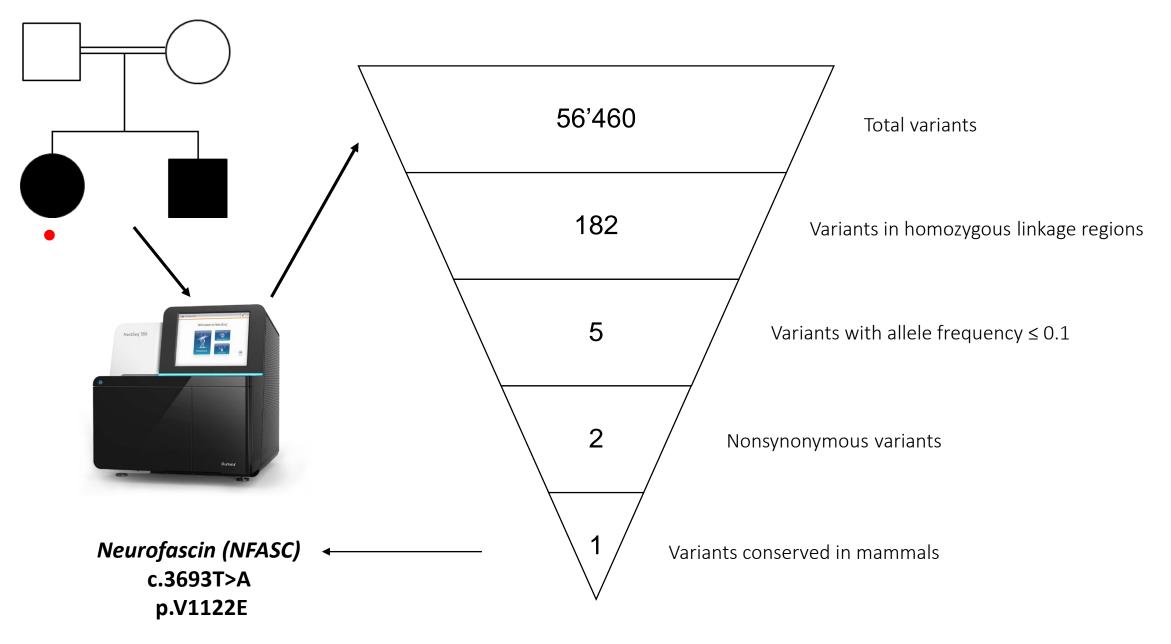


Illumina beadchip SNP-array Genotyping of all core family members

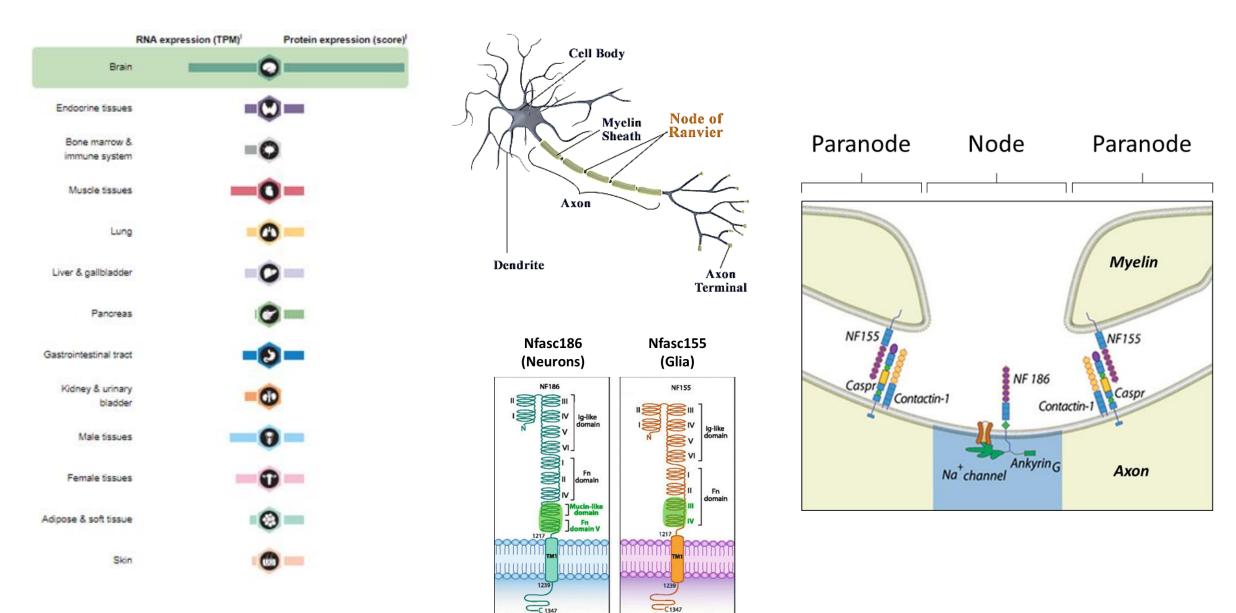


Two homozygous regions shared by the two affected siblings

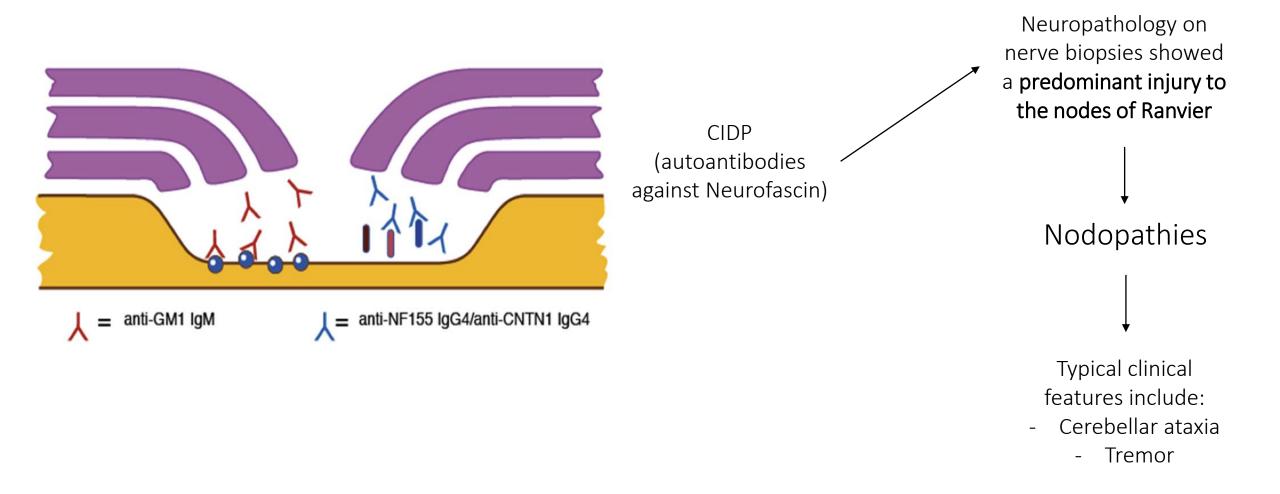
Whole-exome sequencing – filtering analysis



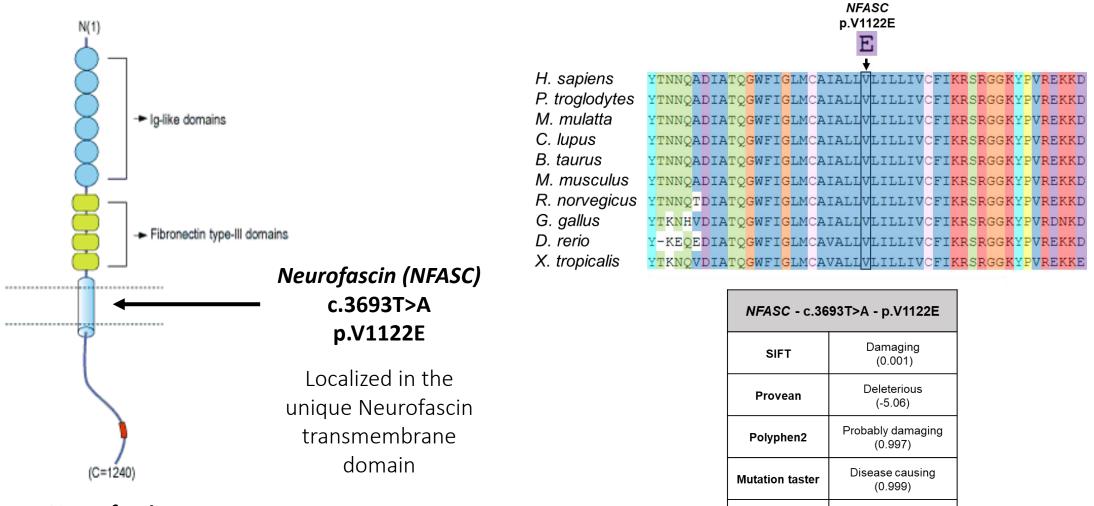
Neurofascin and the node of Ranvier



Neurofascin and human disease (CIDP)



In silico evidence of mutation pathogenicity



Damaging

(0.565)

Deleterious

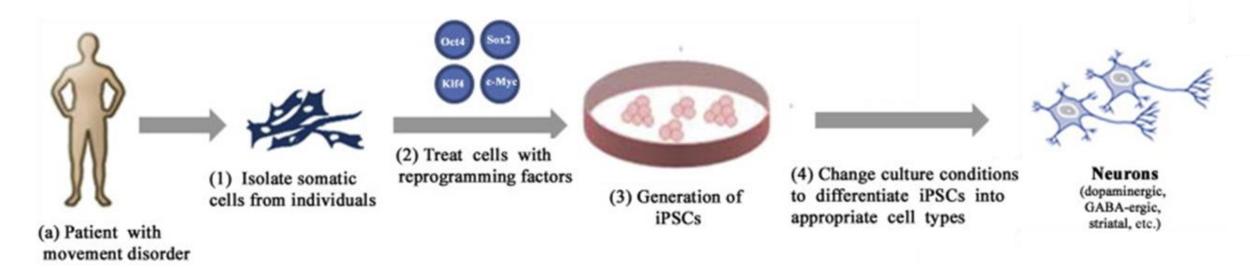
(26.4)

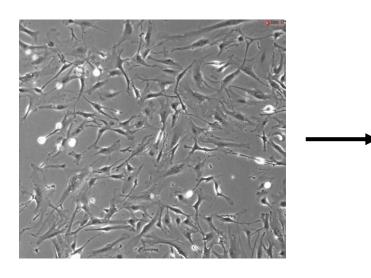
Condel

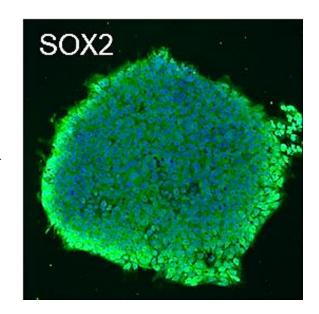
CADD - PHRED

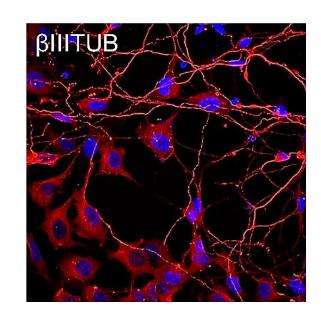
Neurofascin protein domains

In vitro evidence of mutation pathogenicity







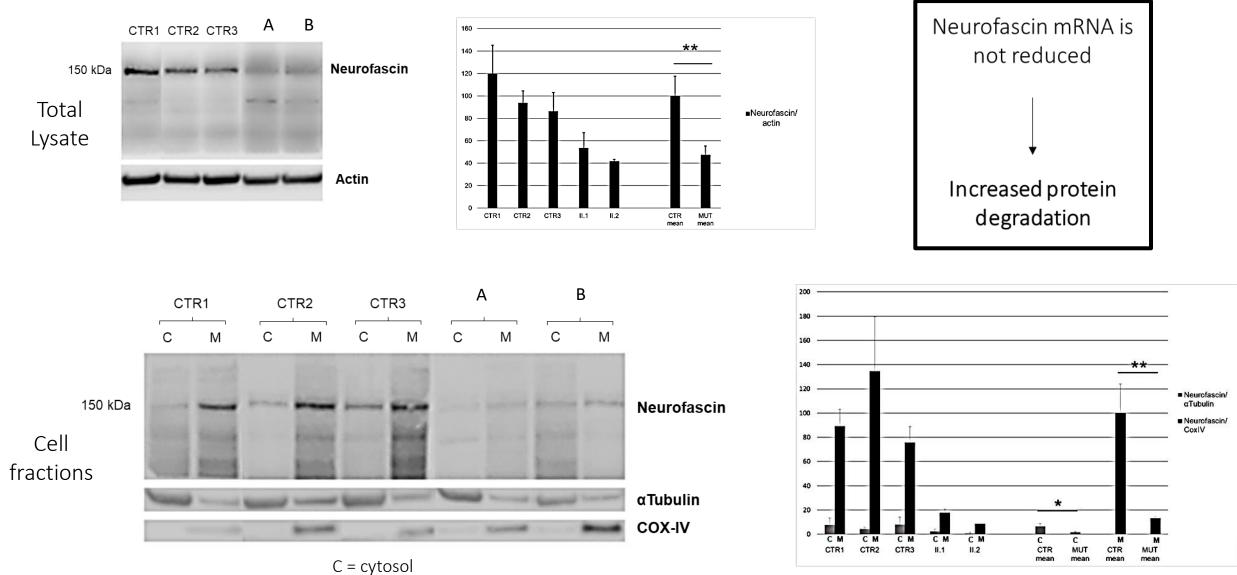


Immunocytofluorescence

Healthy controls Mutated DAPI Neurofascin Merge DAPI Neurofascin Merge A' Neurofascin fluorescence CTR1 * 120 100 80 60 D" D' В D B' 40 20 CTR2 0 m CTR mean MUT mean A В B B'

∢

Western blot (Healthy controls vs patients)



M = membranes

Publications



Parkinsonism & Related Disorders Available online 1 March 2019 In Press, Corrected Proof (?)



Neurofascin (*NFASC*) gene mutation causes autosomal recessive ataxia with demyelinating neuropathy

Edoardo Monfrini ^{a, b}, Letizia Straniero ^{c, d}, Sara Bonato ^{a, b}, Giacomo Monzio Compagnoni ^{a, b}, Andreina Bordoni ^{a, b}, Robertino Dilena ^e, Paola Rinchetti ^{a, b}, Rosamaria Silipigni ^f, Dario Ronchi ^{a, b}, Stefania Corti ^{a, b}, Giacomo P. Comi ^{a, b}, Nereo Bresolin ^{a, b}, Stefano Duga ^{c, d}, Alessio Di Fonzo ^{a, b} 옷 崎

Homozygous mutation in the *Neurofascin* gene affecting the glial isoform of Neurofascin causes severe neurodevelopment disorder with hypotonia, amimia and areflexia ³

Robert Smigiel, Diane L Sherman, Małgorzata Rydzanicz, Anna Walczak, Dorota Mikolajkow, Barbara Krolak-Olejnik, Joanna Kosińska, Piotr Gasperowicz, Anna Biernacka, Piotr Stawinski ... Show more

Human Molecular Genetics, Volume 27, Issue 21, 1 November 2018, Pages 3669–3674, https://doi.org/10.1093/hmg/ddy277 Published: 13 August 2018 Article history ▼

CORRIERE DELLA SERA

La storia di **paziente 2: affetti da una** malattia rarissima

Fratello e sorella, colpiti da una mutazione genetica della coordinazione muscolare. Il loro è l'unico caso conosciuto al mondo Un male per il quale a oggi non esiste cura

di Barbara Gerosa

Conclusions and take-home messages

1. Neurofascin (*NFASC*) mutations are associated with **autosomal recessive hereditary ataxia and demyelinating neuropathy**.

2. **NFASC** gene should be included in **NGS gene-panels** for hereditary ataxias and genetic neuropathies.

3. Neurofascin-related hereditary ataxia with neuropathy is the **first described "genetic nodopathy"**.



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