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Adding evidence to the role of *NEUROG1* in congenital cranial dysinnervation disorders

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Abstract

Congenital cranial dysinnervation disorders (CCDDs) are a heterogeneous group of neurodevelopmental phenotypes caused by a primary disturbance of innervation due to deficient, absent, or misguided cranial nerves. Although some CCDDs genes are known, several clinical phenotypes and their aetiologies remain to be elucidated. We describe a 12-year-old boy with hypotonia, developmental delay, sensorineural hearing loss, and keratoconjunctivitis due to lack of corneal reflex. He had a long expressionless face, severe oromotor dysfunction, bilateral agenesis/severe hypoplasia of the VIII nerve with marked atresia of the internal auditory canals and cochlear labyrinth malformation. Trio-exome sequencing identified a homozygous loss of function variant in the NEUROG1 gene (NM_006161.2: c.202G > T, p.Glu68*). NEUROG1 is considered a causal candidate for CCDDs based on (i) the previous report of a patient with a homozygous gene deletion and developmental delay, deafness due to absent bilateral VIII nerves, and severe oromotor dysfunction; (ii) a second patient with a homozygous NEUROG1 missense variant and corneal opacity, absent corneal reflex and intellectual disability; and (iii) the knockout mouse model phenotype which highly resembles the disorder observed in humans. Our findings support the growing compelling evidence that loss of NEUROG1 leads to a very distinctive disorder of cranial nerves development.

KEYWORDS

aplasia/hypoplasia of sensory cranial ganglia, congenital cranial dysinnervation disorder, *NEUROG1*, oromotor dysfunction, sensorineural deafness, trigeminal nerve aplasia/hypoplasia, vestibulo-cochlear nerve aplasia/hypoplasia

1 | INTRODUCTION

The abnormal development of the cranial nerves can have a genetic basis and result in several phenotypes including recessive hearing loss and congenital cranial dysinnervation disorders (CCDDs), such as Moebius syndrome, Duane syndrome, Marcus Gunn jaw-winking syndrome, congenital ptosis, and congenital fibrosis of extraocular

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muscles.¹ These disorders are heterogeneous and incompletely understood.

NEUROG1 encodes Ngn1 which is a basic helix-loop-helix transcription factor essential for the formation of the proximal cranial sensory ganglia for cranial nerves V and VIII. Ngn1 induces neurogenesis and inhibits the differentiation of neural stem cells into astrocytes by two independent mechanisms. Initially expressed in the epithelium of the trigeminal and otic placodes prior to the onset of neuroblast migration, Ngn1 dimerizes with ubiquitous bHLH proteins and triggers

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a cascade of downstream bHLH factors, including NeuroD, MATH3 and NSCL1, leading to neuronal differentiation. In addition, Ngn1 inhibits astrocyte differentiation by sequestering the CREB-binding protein transcription complex away from astrocyte differentiation genes and by inhibiting the activation of STAT transcription factors necessary for gliogenesis.^{2,3}

We report a patient with mild intellectual disability, epilepsy, deafness, peripheral sensory neuropathy, loss of facial sensation, abnormal development of sensory cranial nerves V and VIII, and a homozygous putative loss of function variant in *NEUROG1*. The patient's phenotype is highly similar to two previously reported cases with homozygous variants in *NEUROG1*.^{4,5} Although *NEUROG1* is not yet associated with a human phenotype in OMIM, our findings support a causal link between *NEUROG1* and a CCDD.

2 | PATIENT AND METHODS

Clinical data was obtained from clinical files and a detailed physical exam was performed. All procedures had the approval of the ethics committee of Hospital de Dona Estefânia, Centro Hospitalar Universitário de Lisboa Central.

Informed written consent was obtained from the patient's parents. All genetic studies were performed for diagnostic purposes.

Clinical data from our patient was compared with published data from two other patients described in the medical literature with homozygous variants in *NEUROG1.*^{4,5}

The patient and his parents were exome sequenced at CENTOGENE AG (Rostock, Germany), as previously described.⁶ Libraries were sequenced on an Illumina Hiseq 4000 with 150 bp paired end sequencing. Variant calling was performed using GATK. Variants with a phred-scaled quality score > 215^7 and a MAF of <1% were retained for analysis. Variants were further filtered based on zygosity and segregation in the family. The database CentoMD®⁸ was queried for rare variants (<1%) in *NEUROG1* (14 August 2020) and the resulting data was filtered taking into consideration an autosomal recessive mode of inheritance and the impact of the variant (non-synonymous).

3 | CLINICAL REPORT

Our patient is a 12-year-old male, the only child of healthy parents of Portuguese origin with no known consanguinity.

Gestation was uneventful and delivery was by caesarean section at term due to feto-pelvic disproportion. Birth weight was 3395 g (50th centile), length 51.5 cm (50–75th centile) and head circumference 34 cm (25–50th centile). Apgar score was 9/10.

He was globally hypotonic since birth. At 22 days he was admitted to hospital due to hypotonia and feeding difficulties requiring naso-gastric feeds.

He had global developmental delay with no history of regression. He sat alone at 18 months and walked independently at 3 years. He never developed expressive language and Brainstem Auditory Evoked Responses at 15 months confirmed profound bilateral sensorineural deafness. He was educated in a school for deaf children and learnt to communicate using sign language.

Ophthalmology review at 18 months was normal; however, at 24 months he had keratoconjuntivitis and lost vision in his right eye. Around the age of five he manifested insomnia, hyperkinesia and selfinjurious behavior (hitting his face and eyes). When he was seven he also presented brief bilateral myoclonic seizures.

At the age of 10 his global IQ (evaluated with appropriate tests for deaf children) was 62, and he manifested major problems with attention, short term memory and abstract thinking.

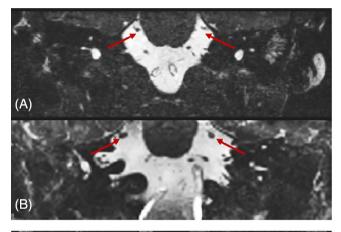
On physical examination at 10 years his stature was 129 cm (10th centile), weight was 24 kg (third centile), and head circumference 52 cm (25th centile). He had a long, asymmetric and expressionless face with open mouth (Figure 1). Long palpebral fissures, high arched eyebrows, ocular leucoma in the right eye, and an arched, and narrow palate were noted (Figure 1). Neurological examination revealed deafness, apparent bilateral loss of facial sensation, and absence of corneal reflexes. Ocular motility was normal. Facial movements in emotional expressions were asymmetric with the mouth deviating to the left. Lower cranial nerve examination was normal. Gait was unsteady with no preponderant side. Muscle power and tone were normal. Stretch reflexes were absent and there was a bilateral flexor plantar response.

Extensive neurometabolic investigations, abdominal ultrasound and echocardiogram were all normal. EMG at 5 years was compatible with an axonal sensory neuropathy. Blink reflexes were absent bilaterally. Brain MRI revealed hypoplasia of cranial nerves V and VIII, marked atresia of the internal auditory canal and hypoplasia of the cochlear labyrinth, bilaterally (Figure 2). The brainstem, cerebellum and cerebral hemispheres were normal. The EEG documented bilateral rolandic spikes with normal background activity.

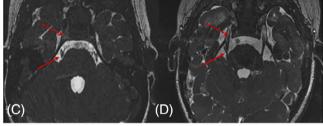
Array CGH showed no relevant copy number variants. Previous molecular testing of DMPK and SLC52A1/SLC52A2/SLC52A3 were also negative. Trio-exome performed for diagnostic purposes (CENTOGENE AG) did not detect relevant variants in morbid genes. An extended search to include genes not previously associated with human disease identified a novel nonsense homozygous variant in NEUROG1 (NM_006161.2): c.202G > T, p.Glu68*. The gene contains only one exon, and there is only one known NEUROG1 protein isoform. The variant introduces a premature termination codon, leading most likely to a truncated protein. Schematic from BAM files showing the location of the variant and the bHLH protein domain is depicted in Figure S1. Both parents were heterozygous for the NEUROG1 variant thus confirming homozygosity in the index case. There are only a few LoF variants in NEUROG1 reported in gnomAD, none in homozygosity (acessed August 14, 2020). Similarly, a search in CENTOGENE data repositories (PMID: 28116331) did not identify additional individuals with homozygous or combined heterozygous LoF variants in this gene.



FIGURE 1 Frontal and profile views of our patient with a homozygous nonsense variant in *NEUROG1* gene at 10 years: (A) long expressionless face, elongated palpebral fissures, facial asymmetry, deviation of the mouth to the left when smiling with loss of the nasolabial fold on the right-side; (B) inability to close the mouth [Colour figure can be viewed at wileyonlinelibrary.com]



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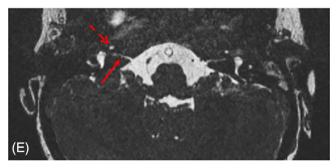


FIGURE 2 T2 weighted brain MRI, Constructive Interference in Steady State (CISS) sequence: (A),(B) Coronal view of the patient's trigeminal nerves (A) showing they are much thinner than in a normal control at the same age (B); (C),(D) Axial view of the patient's trigeminal nerves (C) in comparison with a normal control (D) with the dashed arrows pointing at the trigeminal ganglion; (E) Axial view showing a stenotic internal auditory canal (arrow) and hypoplastic cochlea (dashed arrow) [Colour figure can be viewed at wileyonlinelibrary.com]

4 | DISCUSSION

Several genes related to neuronal development are currently linked to congenital cranial dysinnervation disorders (*PLXND1* and *REV3L* for Moebius syndrome, *MAFB*, *HOXA1*, *SALL4* and *CHN1* for Duane syndrome, *ROBO3* for horizontal gaze palsy with progressive scoliosis, and *PHOX2A*, *KIF21A*, *TUBB3* and *TUBB2B* for congenital fibrosis of extraocular muscles). These disorders are believed to result from a primary disruption of innervation due to the abnormal development of cranial nerves.⁹⁻¹³

In the developing CNS the formation of trigeminal and vestibulocochlear nerves requires the activity of *NEUROG1*.^{2,3} The development of dorsal root ganglion cells is likewise dependent on *NEUROG1*, which also commits neuronal progenitor cells in the olfactory bulb, cerebral cortex, hippocampus, thalamus, brainstem and cerebellum.¹⁴ The gene *NEUROG1* and protein Ngn1 have been well studied in vertebrate animal models.^{15,16}

Ma et al. (1998) generated Ngn1-null mice embryos showing complete absence of trigeminal and otic placode-derived vestibulocochlear ganglia, apparent truncation of cranial nerve V, and loss of *expression of* Ngn1 *downstream bHLH factors, including NeuroD, Math3 and* NSCL1, *in mutant otic and trigeminal placodes.*

Present knowledge about the biology of Ngn1 and the phenotype of the animal model are concordant with the clinical features we report here. Our patient's deafness, hypoplastic cranial nerve VIII, bilateral atresia of the internal auditory canals, and hypoplasia of the cochlear labyrinth match the observations in Neurog1 knock-out mice.^{15,16} Ngn1-null mutants had no vestibulocochlear ganglion, no VIIIth nerve afferent, efferent and autonomic fibres, small abnormal inner ear and cochlear hypoplasia. Our patient's hypoplasia of cranial nerve V, lack of corneal reflexes, and facial anaesthesia were associated with oromotor dysfunction and led to a corneal traumatic lesion. In mice lacking Ngn1 inability to suckle caused early lethality. In the human patients medical resources such as parenteral feeding, nasogastric tube and interventions focusing on strengthening oromotor function helped overcome equivalent feeding difficulties. *NEUROG1* is involved in neurogenesis in the central nervous system including the neocortex.¹⁷ Although the intellectual disability present in our patient may be partly ascribed to deafness and sensorial deprivation, and the self-injurious behavior likewise attributed to abnormal sensation in the trigeminal territory, it is likely these features, as well as epilepsy, are manifestations of abnormal cortical development. *NEUROG1* promotes sensory neuron differentiation in dorsal root ganglia.¹⁸ Thus, delayed motor development, and the findings on nerve conduction studies compatible with a subclinical axonal sensory neuropathy in our patient, may also be related to the *NEUROG1* gene defect.

In 2013, Schroder et al.⁴ reported one patient with dysmorphic features, mild intellectual disability, deafness, severe hypoplasia of the vestibulo-cochlear nerves, narrow internal auditory canal, hypoplastic cochlea, and oromotor dysfunction. This patient had a homozygous microdeletion in chromosome 5 including all exons and corresponding promoter regions of the *NEUROG1*, *TIFAB* and *DCNP1* genes. The authors claimed the phenotype in Neurog1 knock-out mice was in agreement with the clinical and radiological findings in their patient. Moreover, they observed a balance disorder in their patient, which

they linked to the absence of utriculosaccular ducts in Ngn1-null mice. Malfunction of cranial nerve V was assumed, but loss of sensation in this nerve's territory or its absence on brain MRI were not described. Based on the reported phenotype of Ngn1-null mutant mice, Schroder et al. proposed *NEUROG1* as a new candidate gene for a congenital cranial dysinnervation disorder including cranial nerves V and VIII, and excluded *TIFAB* and *DCNP1* as causative.¹⁹⁻²¹

Finally, in 2015, Yavarna et al. identified a patient with corneal opacity, absent corneal reflexes and intellectual disability who had a homozygous missense variant in *NEUROG1*: c.347G > T, p. Arg116Leu.⁵ A summary of the clinical phenotypes of these patients is shown in Table 1.

Taken together, the very unusual combination of clinical features, and the striking similarity in findings between these three patients and the characteristics of the knock-out mouse model, provide compelling evidence that loss of NEUROG1 is causing CCDDs in humans. Our data strengthens the association of *NEUROG1* with a very distinctive phenotype that comprises features related to hypoplasia/absence of the Vth and VIIIth cranial nerves, mild intellectual disability and, possibly, a wider clinical spectrum including manifestations such as

TABLE 1	Comparison of clinica	I data between Schroder and	Yavarnas's cases and our patient
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	Schroder et al., 2013	Yavarna et al., 2015	Our patient
Neonatal period			
Normal birth parameters	Yes	nk	Yes
Poor sucking and swallowing	nk	nk	Yes
Global hypotonia	Yes	nk	Yes
Torticollis	Yes	nk	No
Growth and development			
Weight and height between 3rd-97th centiles	Yes (weight 50th centile and height 5th centile)	nk	Yes (weight 3th centile and height 10th centile)
Head circumference between 3rd-97th centiles	Yes (75th centile)	nk	Yes (25th centile)
DD/ID	Yes (mild)	Yes (mild)	Yes (mild)
Neurological involvement			
Lagopthalmos, absent corneal reflex	nk	Yes	Yes
Bilateral loss of facial sensation	nk	nk	Yes
Bilateral sensorineural hearing loss	Yes (profound)	nk	Yes (profound)
Oromotor dysfunction	Yes (increased salivation, inability to chew, mashed diet)	nk	Yes (chewing/swallowing difficulties, unable to close the mouth)
Balance disorder	Yes	nk	Yes (ataxic gait)
Muscle weakness	Yes	nk	No
Epilepsy	No	nk	Yes (brief bilateral myoclonic seizures)
Craniofacial features			
Cranial conformation	plagioscaphocephaly	nk	plagiocephaly
Long face	Yes	nk	Yes
Lack of facial mimicry	Yes	nk	Yes
Facial asymmetry	Yes	nk	Yes
High arched eyebrows and elongated palpebral fissures	Yes	nk	Yes

TABLE 1 (Continued)

	Schroder et al., 2013	Yavarna et al., 2015	Our patient
Low-set posteriorly rotated ears	Yes	nk	No
High and narrow palate	Yes	nk	Yes
Other features			
Pectus excavatum	Yes	nk	No
Sacral dimple/ hypoplastic genitalia	Yes	nk	No
Faulty foot posture	Yes	nk	No
Neuroimaging findings			
Bilateral agenesis of cranial nerve VIII	Yes	nk	Yes
Bilateral agenesis of cranial nerve V	nk	nk	Yes
Atresia/stenosis of the internal auditory canals	Yes	nk	Yes
Cochlear labyrinth malformation	Yes	nk	Yes
Other neurological exams			
Electromyography	nk	nk	Axonal sensory neuropathy
Electroencephalogram	nk	nk	Bilateral rolandic spikes with a normal background activity
Genetic findings			
Genetic testing results	Homozygous deletion of 116.796 kb at 5q31.1, involving NEUROG1	Homozygous missense variant in NEUROG1: c.347G > T, p.Arg116Leu	Homozygous frameshift variant in NEUROG1 (NM_006161.2): c.202G > T, p.Glu68*

Abbreviations: DD/ID, developmental delay/intellectual disability; nk, not known.

epilepsy and axonal sensory neuropathy; and contributes to a better understanding and classification of these disorders.

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CONFLICT OF INTEREST

The authors declare that they have no conflict of interest.

PEER REVIEW

The peer review history for this article is available at https://publons. com/publon/10.1111/cge.13922.

DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available on request from the corresponding author. The data are not publicly available due to privacy or ethical restrictions.

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of this article.

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