

## RESEARCH ARTICLE

# International Registry of *NKX2-1*-Related Disorders: Clinical, Genetic, and Imaging Perspectives

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**ABSTRACT: Background:** *NKX2-1*-related disorders result from heterozygous variants in *NKX2-1*, a gene crucial for brain, lung, and thyroid development. Although movement disorders, hypothyroidism, and neonatal respiratory distress are recognized, the full phenotype and genotype-phenotype relationships remain incompletely defined.

**Objectives:** To delineate neurological, respiratory, and endocrine features across ages, characterize movement disorder trajectories – particularly chorea – and explore genotype-phenotype associations with clinical relevance.

**Methods:** We conducted a multicenter, cross-sectional study recruiting participants through referral clinicians and European networks. Standardized clinical and genetic data were captured in an electronic database and analyzed with descriptive and inferential statistics.

**Results:** Sixty-eight individuals (37 female; median age 16 years, range 2–60 years) were included. Motor delay was the commonest presenting feature (~60%); neonatal respiratory distress syndrome occurred in one-third of cases. The brain-lung-thyroid triad was present in almost half. Chorea affected over 90% and began in early childhood; it was more frequent with single nucleotide variants than with deletions. Deletions are

associated with better gross motor function. Frameshift or nonsense variants showed greater respiratory involvement, and variants in the exon-3 homeobox region were associated with age-related reduction of chorea. Neonatal respiratory distress predicted later respiratory symptoms. Greater abnormal involuntary movement severity correlated with poorer manual and gross motor function. Hypotonia and untreated hypothyroidism are associated with more severe chorea. Psychiatric comorbidity occurred in over one-third of cases, mainly attention-deficit/hyperactivity symptoms.

**Conclusions:** This largest cohort to date shows early neurological onset, genotype-specific outcomes, and frequent psychiatric comorbidity in *NKX2-1*-related disorders, refining clinical expectations and supporting genotype-informed diagnosis, counseling, and management. © 2026 International Parkinson and Movement Disorder Society.

**Key Words:** chorea; benign hereditary chorea; brain-lung-thyroid syndrome; hypothyroidism; neonatal respiratory distress syndrome; neurodevelopmental delay; *NKX2-1*; TTF-1

*NKX2-1*-related disorders (*NKX2-1*-RD), also known as brain-lung-thyroid syndrome or benign hereditary chorea, represent a clinically and genetically diverse group of conditions caused predominantly by pathogenic single nucleotide variants (SNVs) in the *NKX2-1* gene, located on chromosome 14q13.<sup>1,2</sup> The gene encodes the thyroid transcription factor-1 (TTF-1), a homeodomain-containing transcription factor essential for embryonic development and postnatal function of the brain, lungs, and thyroid gland. *NKX2-1*-RD typically exhibits an autosomal dominant inheritance, although *de novo* cases have also been reported.

The classic presentation of *NKX2-1*-RD is characterized by a triad of features: movement disorders (most commonly chorea, though dystonia, myoclonus, and ataxia may also occur), thyroid dysfunction (ranging from congenital hypothyroidism to compensated forms), and pulmonary involvement, often manifesting as neonatal respiratory distress syndrome (NRDS).<sup>1</sup> The phenotypic spectrum may sometimes include one or two components of the triad or extend beyond it to additional neurological, respiratory, and endocrine manifestations. There is also emerging evidence for increased predisposition to thyroid and lung malignancies and for psychiatric comorbidities.<sup>3,4</sup>

The *NKX2-1* protein regulates genes essential for thyroid hormone synthesis (eg, thyroglobulin, thyroperoxidase, and thyrotropin receptor),<sup>5–7</sup> supports differentiation of alveolar type II cells and surfactant production in the lungs,<sup>8–11</sup> and contributes to the development of the basal

ganglia, hypothalamus, and other structures involved in motor control and neuroendocrine regulation.<sup>7</sup> Disruption of these functions results in multisystem disease with variable expressivity and penetrance, even among individuals with identical variants.

Despite the role of *NKX2-1* in these disorders being established over two decades ago, important gaps remain in understanding the complete clinical spectrum, natural history, and genotype-phenotype correlations.<sup>12,13</sup> Current knowledge is largely derived from case reports and small cohorts, limiting insight into phenotypic diversity.<sup>1,14–30</sup> For example, while chorea is a hallmark neurological feature, its longitudinal course is poorly defined<sup>31</sup>; the longest reported follow-up spans 24.5 years and includes 28 individuals.<sup>29</sup> The frequency and severity of respiratory and thyroid manifestations also vary, and correlations with genotype are only beginning to emerge.<sup>32,33</sup> Recent studies have expanded the genetic spectrum to include deletions, mobile element insertions, changes in conserved non-coding regions, and variants in regulatory genes (eg, *PAX9*, *MBIP*) that may influence phenotype.<sup>34,35</sup>

This heterogeneity complicates diagnosis and management. Symptom overlap with other neurodevelopmental, movement, and endocrine disorders often delays diagnosis. In addition, the unpredictable evolution of symptoms makes prognosis and long-term treatment planning challenging.

To address these gaps, this study analyzed the largest multicenter cohort of genetically confirmed *NKX2-1*-RD

to date. Through detailed clinical, genetic, and neuroimaging assessments, we aimed to define the full range of neurological, respiratory, and endocrine manifestations across ages, characterize the progression of movement disorders – particularly chorea – and identify genotype–phenotype associations that can guide diagnosis, prognostication, and targeted management strategies.

## Patients and Methods

### Study Design and Population

This multicenter, cross-sectional, observational study included individuals with NKX2-1-RD and a confirmed genetic diagnosis. Participants were recruited through referral physicians and international outreach via several platforms, including the European Reference Network for Rare Neurological Diseases (ERN-RND), the European Reference Network for Rare Malformation Syndromes, Intellectual and Other Neurodevelopmental Disorders (ERN-ITHACA), the Spanish Society of Pediatric Neurology (SENEP), the Pediatric Movement Disorders Special Interest Group (SIG) of the International Parkinson and Movement Disorder Society (MDS), and the Facebook NKX2-1 patient group. Additional recruitment was achieved through workshops, scientific congresses, and direct referrals from colleagues.

All study data were collected and managed in a RED-Cap (Research Electronic Data Capture) database for the International NKX2-1 Registry,<sup>36,37</sup> hosted at Hospital Sant Joan de Déu, Barcelona, Spain, between October 2023 and October 2024. In total, 40 specialists from 29 centers across 17 countries contributed to the registry. Clinical data were obtained from medical records and de-identified to ensure confidentiality.

Clinical and ancillary data were collected using a standardized registry protocol and analyzed following predefined criteria. Core variables included demographic information, perinatal history, neurological findings, systemic features, and genetic results. Movement-disorder phenomenology was assessed by subspecialists at each site, and severity was graded using the Abnormal Involuntary Movement Scale (AIMS) when available. Brain magnetic resonance imaging (MRI) and endocrine assessments were reviewed locally. Genetic testing followed site-specific protocols and American College of Medical Genetics and Genomics (ACMG)/Association for Molecular Pathology (AMP) classification guidelines (NKX2-1, transcript NM\_001079668.3). Statistical analyses included descriptive, univariate, and multivariate models using appropriate nonparametric tests (Spearman, Mann–Whitney, Kruskal–Wallis, Chi-squared/Fisher) with  $P < 0.05$  considered significant.

A detailed description of data collection procedures, operational definitions (eg, gait abnormalities, cognitive

impairment), and statistical methods is provided in Supplementary Material, Methods in Data S1.

## Results

### Demographic and Neurological Features

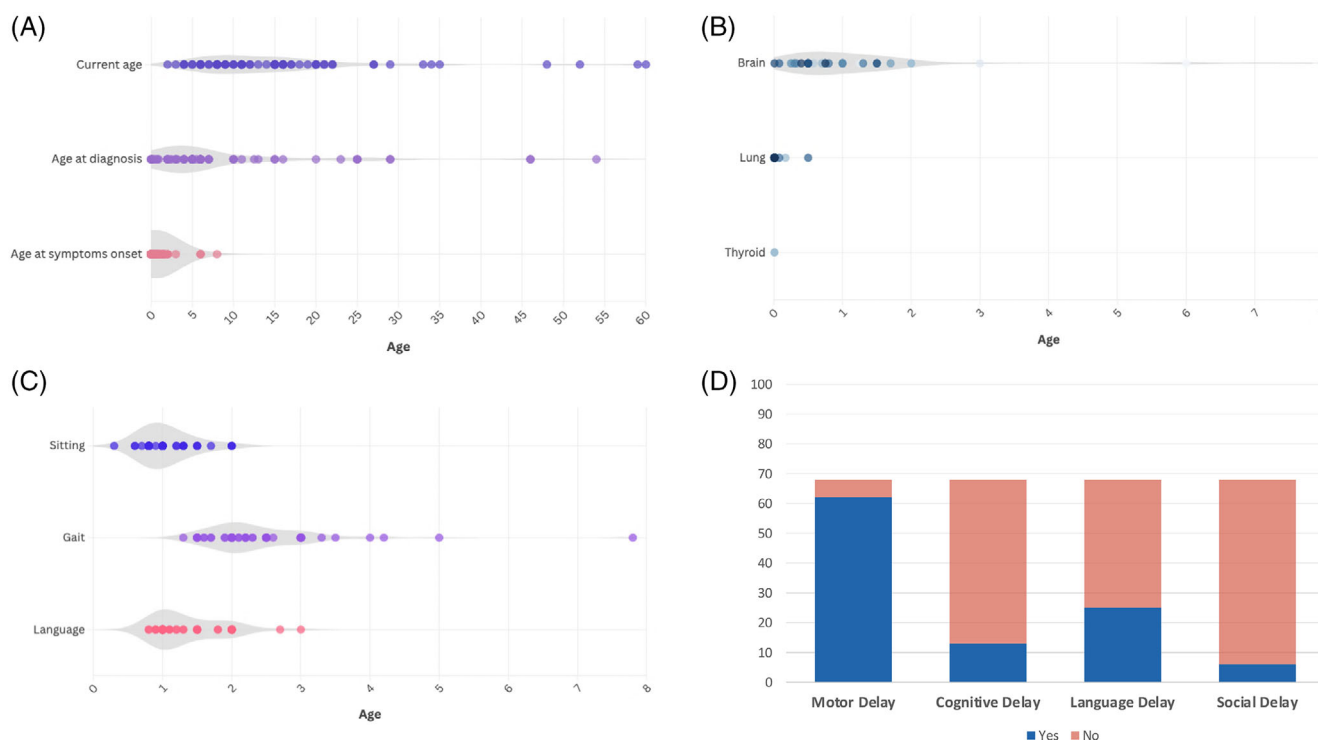
Sixty-eight individuals with NKX2-1-RD were included; 37 (54.4%) were female. Median age at last follow-up was 16 years (range 2–60 years) (Fig. 1A). Most were referred by neurologists (86.8%). Eighty-five percent were followed in Europe (notably France and Spain) and 15% were in the Americas. One neonatal death due to NRDS occurred. Fourteen cases had been previously published.<sup>5,22,28,29,38–43</sup>

Median age at first symptom was within the first year (median 0.87 years, interquartile range [IQR] 0–8 years) (Fig. 1A). Neurological symptoms predominated at onset: motor delay was most frequent (41.8%), followed by global developmental delay and hypotonia; chorea or gait problems were initial in a minority. Respiratory onset occurred in 34.3% (mostly NRDS) and endocrine onset was rare. Only half (50.8%) fulfilled the full brain–lung–thyroid triad; 40% had dual-system involvement (most often brain–thyroid), and 9.2% showed isolated brain involvement (Fig. 1B). A phenotypic heatmap for age and sex is shown in Figure 2.

Perinatal complications were common. NRDS was reported in 35.5% (median onset 4.5 hours after birth). About half of those with NRDS required invasive ventilation; others received noninvasive support or oxygen. Additional neonatal issues (25.81%) included hyperbilirubinemia, non-reassuring fetal status (eg, abnormal fetal heart rate patterns), and late prematurity. Among term infants, median birthweight was 3135 g; newborn screening was abnormal in 22.7%, all subsequently diagnosed with hypothyroidism (Supplementary Table S1).

Neurodevelopmental delay (NDD) was present in 94% (64/68): mild 62.1%, moderate 31.0%, and severe 6.9% (when graded). Motor delay was nearly universal (91.2%), whereas speech (36.8%), other cognitive (19.1%), and social (8.8%) impairments were less frequent (Fig. 1C,D). Independent sitting occurred at 12.4 months on average; autonomous gait at 29.5 months. Only 10.7% walked before 19 months, and later gait onset correlated with greater NDD severity (Spearman  $r = 0.524$ ,  $P < 0.001$ ) (Fig. 3A).

Chorea affected 92.7% (Supplementary Table S2). It began early (median 2.00 years; 93% before age 6 years). Chorea was predominantly generalized (80.6%). Among patients with non-generalized chorea, the most frequently affected body regions were the arms (66.7%), head (22.2%), and legs (11.1%). Chorea



**FIG. 1.** Clinical characterization of individuals with NKX2-1-related disorders (NKX2-1-RD). (A) Violin plot depicting current age (in years), age at NKX2-1-RD diagnosis, and age at onset of first symptoms. (B) Violin plot showing the age (in years) at onset of brain, lung, and/or thyroid symptoms. (C) Violin plot illustrating the age (in years) at attainment of sitting, gait, and language milestones. (D) Column chart showing the percentage of individuals with NKX2-1-RD presenting motor, cognitive, language, and social developmental delays. [Color figure can be viewed at [wileyonlinelibrary.com](http://wileyonlinelibrary.com)]

was more frequent with NKX2-1 SNVs than with deletions (96.6% vs. 75%, Fisher's exact test  $P = 0.067$ ). Females showed earlier onset when univariate Mann-Whitney U test analysis was performed, but this information was not retained after adjustment in the multivariable linear regression.

Evolution of chorea could be classified in those with  $\geq 2$  documented assessments; given the heterogeneous and non-uniform observation intervals across patients, results are reported as categorical evolution (improved/stable/worsened) rather than as time-to-change. Over time, chorea stabilized in 57.1%, improved in 35.7%, and worsened in 7.1%. Females appeared to have better outcomes in unadjusted analysis (Fisher's exact test  $P = 0.035$ ) (Fig. 3C), but this did not persist in multivariable logistic regression controlled by possible confounders. Two clinical factors associated with a worse chorea course were hypotonia (Fisher's exact test  $P = 0.002$ ) and untreated hypothyroidism (Fisher's exact test  $P = 0.032$ ) (Fig. 3D,E). By genotype, homeobox exon-3 SNV tended to remain stable, whereas non-homeobox SNV more often improved (Fisher's exact test  $P = 0.039$ ) (Fig. 3F). On the AIMS rating scale ( $n = 52$ ), 25% had minimal, 57.7% mild, and 17.3% moderate chorea.

Other motor features were frequent (82.4%): hypotonia 57.4%, dystonia 46.3%, gait abnormalities

34.3%, myoclonus 32.8%, dysarthria 14.9%, and tremor 9.0%. Frequent falls were reported in 61% of those queried. Myoclonus was associated with dystonia (51.6% vs. 17.1%,  $\chi^2 = 8.790$ ,  $df = 2$ ,  $P = 0.007$ ) and with gait abnormalities/dysarthria (70% vs. 28.6%,  $\chi^2 = 7.299$ ,  $df = 2$ ,  $P = 0.026$ ). Myoclonus was uncommon before the second decade but present in ~49% thereafter ( $\chi^2 = 6.271$ ,  $df = 2$ ,  $P < 0.05$ ) (Fig. 3B). Intellectual disability was documented in 11.5%; most others were in the borderline-mild range.

Functional scales showed mild impairment overall (Gross Motor Function Classification System [GMFCS] I-II in 95%; Manual Ability Classification System [MACS] I-II in 88%). Deletions were uniformly associated with better gross motor function (GMFCS I) than SNV (Fisher's exact test  $P = 0.045$ ). Males had worse manual ability (higher MACS scores) than females (Fisher's exact test  $P = 0.047$ ).

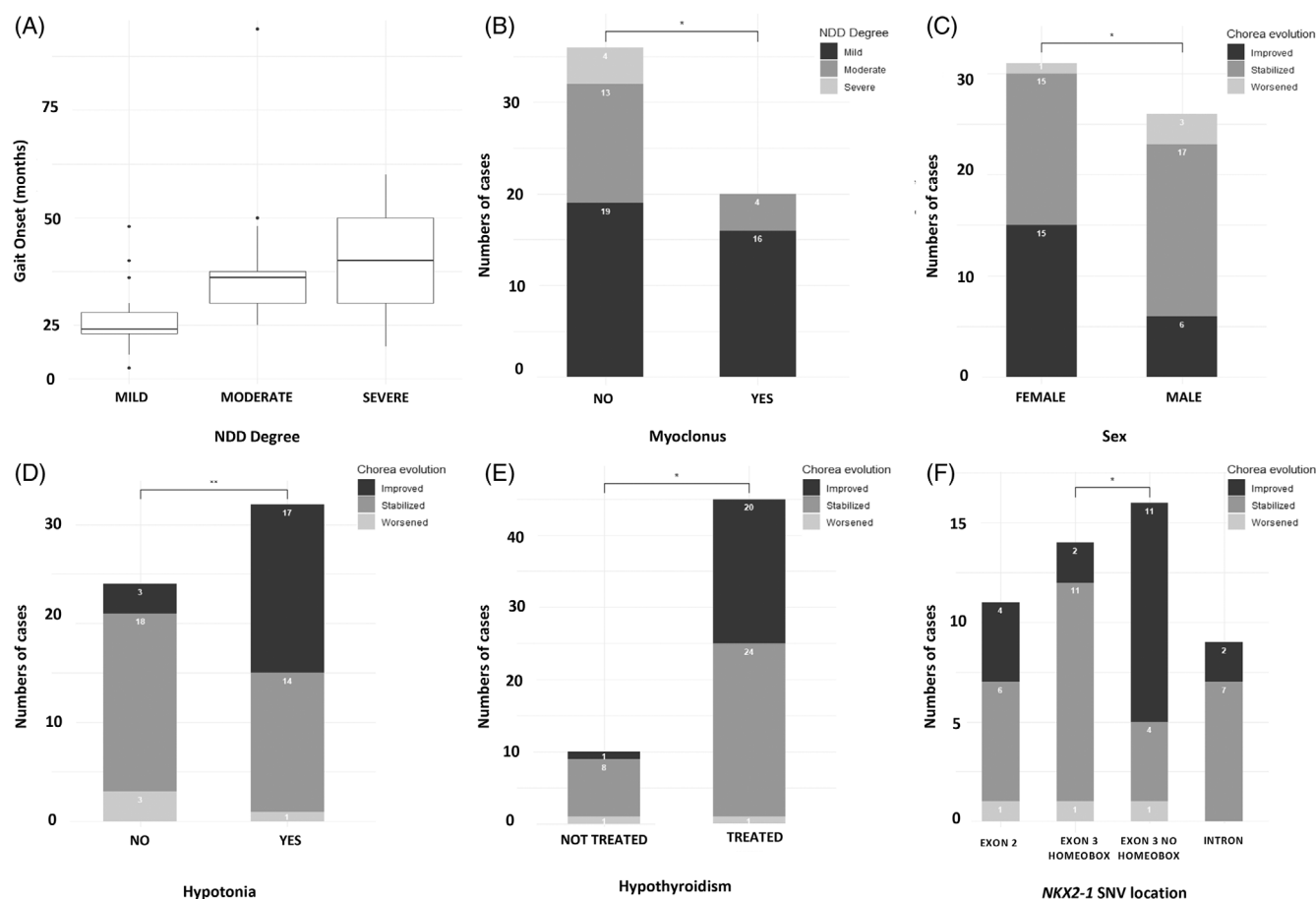
### Systemic Features

Respiratory symptoms occurred in 57.6%; recurrent wheezing and bronchospasm consistent with reactive airway disease were most common (42.9%). NRDS predicted later respiratory disease (87% vs. 45%,  $\chi^2 = 13.816$ ,  $df = 2$ ,  $P = 0.001$ ). The



	AGE (years)	SEX	BRAIN	LUNG	THYROID	CHOREA	NDD	ID	HYPOTONIA	GAIT ABNORMALITIES	DYSTONIA	DYSARTHRIA	MYOCLONUS	TREMOR	FALLS	SEIZURES	ADHD	ANXIETY	DEPRESSION	NRDS	ASTHMA	PULMONAR FIBROSIS	OTHER LUNG SYMPTOMS	HYPOTHYROIDISM	FAILURE TO THRIVE	OTHER ENDOCRINOLOGICAL
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2	59	M																								
3	52	M																								
4	48	F																								
5	35	F																								
6	34	M																								
7	33	F																								
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FIG. 2. Legend on next page.



**FIG. 3.** Clinical associations in individuals with *NKX2-1*-related disorders (*NKX2-1*-RD), focusing on neurodevelopmental delay (NDD) and chorea. (A) Age at gait onset by NDD severity. Boxplot illustrating gait onset age (in months) in individuals with mild, moderate, and severe NDD. Median, interquartile range, and outliers are shown, indicating a significant correlation between delayed gait onset and NDD severity. (B) Distribution of NDD severity in individuals with and without myoclonus. Bar chart showing the number of cases with mild, moderate, or severe NDD, stratified by presence (Yes) or absence (No) of myoclonus. A significant difference in NDD severity was observed between groups. (C) Chorea evolution by sex. Boxplot showing age at chorea onset (months) by sex, with a trend toward earlier onset in females that was not significant in multivariable models. (D) Chorea evolution by presence of hypotonia. Stacked bar chart comparing outcomes of chorea (improved, stabilized, worsened) in individuals with and without hypotonia. A significant difference was observed. (E) Chorea evolution by presence of treated hypothyroidism. Stacked bar chart showing chorea evolution in individuals with and without treated hypothyroidism. A significant difference was observed. (F) Chorea evolution by location of *NKX2-1* single nucleotide variants (SNV). Stacked bar chart comparing chorea evolution across individuals with variants in different gene regions. A significant difference was observed. Statistical significance: \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .

average age at the onset of recurrent wheezing and bronchospasm consistent with reactive airway disease was  $\sim 1.2$  years. Recurrent infections were reported in seven individuals; interstitial lung disease in two; and one adult had unclassifiable pulmonary fibrosis. Only about one-fifth underwent spirometry or chest computed tomography; diffusion capacity was rarely assessed. Treatments included inhaled steroids, bronchodilators, and azithromycin. Frameshift/nonsense SNVs were associated with higher respiratory involvement than other SNVs ( $\chi^2 = 8.270$ ,  $df = 2$ ,  $P = 0.016$ ).

Hypothyroidism was highly prevalent (82%; median diagnosis age was 1.54 years, nearly half in infancy). Median thyroid-stimulating hormone (TSH) at diagnosis was elevated, and median free thyroxine (T4) was low-normal. Thyroid ultrasound (available in half the cases) showed congenital hypoplasia in about one-third. Most hypothyroid individuals (85%) received levothyroxine, with heterogeneous dosing; precise data distinguishing congenital versus compensated cases were incomplete.

Beyond the triad, 19.7% had other endocrine issues: growth hormone deficiency (9.1%), hypogonadotropic

**FIG. 2.** Phenotypic heatmap of individuals with *NKX2-1*-related disorders (*NKX2-1*-RD). The heatmap illustrates the presence or absence of the classical brain–lung–thyroid triad, together with various neurological, respiratory, and endocrinological symptoms, categorized by age and sex. Red indicates absence, blue indicates presence, and white represents missing data. NDD, neurodevelopmental delay; ID, intellectual disability; NRDS, neonatal respiratory distress syndrome; ADHD, attention-deficit hyperactivity disorder; F, female; M, male. [Color figure can be viewed at [wileyonlinelibrary.com](https://onlinelibrary.wiley.com/doi/10.1002/mds.20187)]

hypogonadism (4.6%), and isolated cases of hypopituitarism or early puberty. Anthropometrics were generally within reference ranges, and thyroid dysfunction did not associate with failure to thrive or short stature.

Psychiatric comorbidity was common (over one-third). Attention-deficit/hyperactivity symptoms were most frequent, followed by anxiety and depression. Myoclonus was significantly associated with anxiety and depression ( $\chi^2 = 7.967$ ,  $df = 2$ ,  $P < 0.01$  and  $\chi^2 = 12.776$ ,  $df = 2$ ,  $P < 0.01$ , respectively). No cases of autism spectrum disorder or obsessive-compulsive disorder were recorded. Treatments included methylphenidate (often continued at the last follow-up), guanfacine, or atomoxetine in isolated cases, and selective serotonin reuptake inhibitors or antipsychotics when indicated.

About 31% reported additional non-triad features (eg, fatigue, urinary incontinence, joint hyperlaxity, mild humoral immunodeficiency); oncological processes were not observed. Review of 13 facial photographs did not reveal a consistent dysmorphic pattern.

### Genetic Diagnosis and Neuroimaging

Median age at genetic diagnosis was 5 years. Testing modalities included exome sequencing (clinical or whole; 79%), microarray-based comparative genomic hybridization (8%), whole-genome sequencing (3%), and others. Most individuals (83.8%) harbored *NKX2-1* SNV; the remainder had deletions or regulatory-region disruptions (eg, *MBIP*, *PAX9*). Among SNV, nonsense (40.4%) and frameshift (28.1%) were most frequent, followed by splicing (15.8%) and missense (17.8%). SNVs are distributed across exon 2, exon 3, and splice regions; all missense variants are clustered within the homeobox domain. Most nonsense variants also lay within the homeobox; only one frameshift was located there (distal). Two related individuals had an Alu retrotransposition in exon 3. Eight individuals (11.8%) had deletions (two megadeletions and six microdeletions); five encompassed *NKX2-1* with other genes, and three spared *NKX2-1* but included regulatory partners (*MBIP* in two and *PAX9* in one). One individual had a translocation involving the *NKX2-1* region (Supplementary Fig. S1, Table S3).

According to ACMG, 57.8% of SNVs/copy number variants (CNVs) were pathogenic, 40.6% likely pathogenic, and 1.6% variant of unknown significance (VOUS). One case (I7) harbored a heterozygous 14q13.3 microdeletion (arr[GRCh37] 14q13.3(36,722,498–36,790,795)x1; 68.3 kb) encompassing *MBIP*. The clinical presentation was concordant with *NKX2-1*-RD. According to the ClinGen CNV rubric (evidence codes 1A, 3A, 4N), this CNV remains classified as a VOUS trending toward likely pathogenic at the time of resubmission. Approximately half of the variants were *de novo*; among inherited variants with known parents,

maternal transmission predominated. Seven multiplex families are represented, though parental data were incomplete in most, limiting analyses of penetrance and intrafamilial variability.

Brain MRI (performed in 80.6%) was normal in 67.9%. Reported abnormalities included corpus callosum dysgenesis, delayed myelination, and cystic lesions (7.6%), such as Rathke cleft cysts and an arachnoid cyst. Other findings were empty sella, Chiari I malformation, and non-specific white matter changes.

### Treatment and Management

Movement disorder-directed therapies were used in 35/63 (55.6%). The most frequently prescribed agents were tetrabenazine (54.3% of treated patients), levodopa (45.7%), and methylphenidate (22.9%). Tetrabenazine was initiated at a median age of 6.5 years and was often discontinued due to limited benefit or adverse effects. Levodopa and methylphenidate provided moderate benefit in a subset of patients. At the last follow-up, approximately half of the treated individuals remained on monotherapy. One patient with dystonia underwent globus pallidus internus (GPi) deep brain stimulation with a favorable response (I7). Details are provided in Supplementary Table S2 and Figure 4.

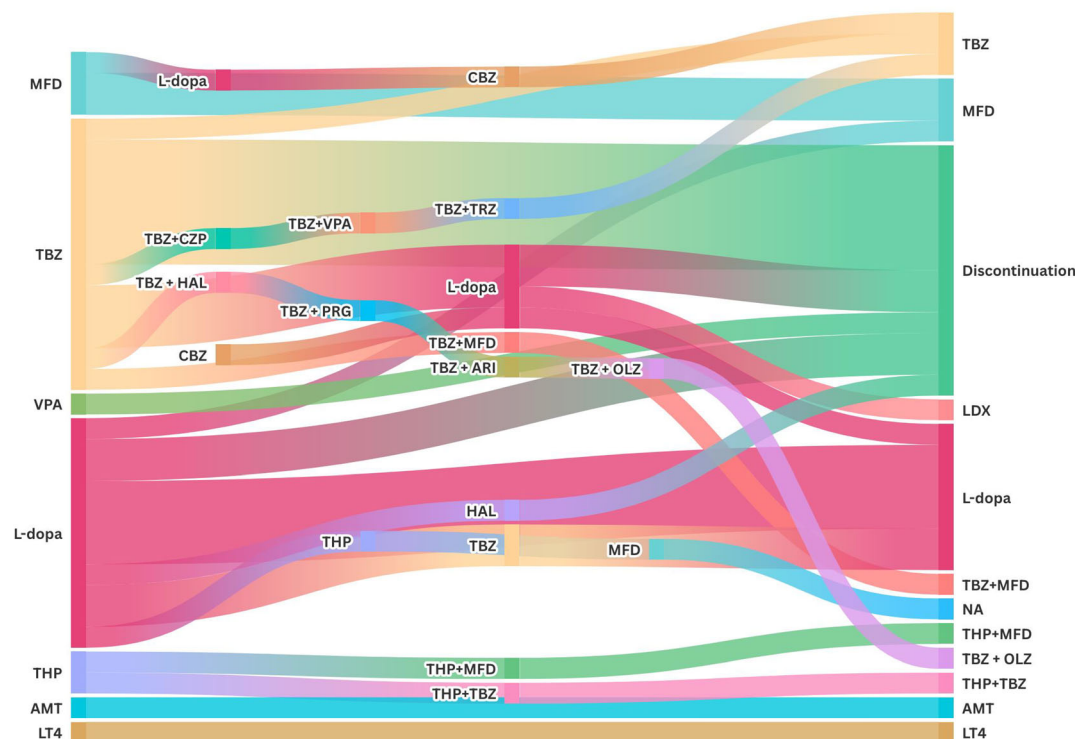
Care was multidisciplinary (neurology, pulmonology, and endocrinology), with periodic assessment of motor function, respiratory status, and thyroid hormones, alongside neurodevelopmental and psychiatric monitoring (Supplementary Table S4).

A more detailed description of the results can be found in the Supplementary Material in Data S1.

### Discussion

Our findings broaden the clinical and genetic understanding of *NKX2-1*-RD and challenge reliance on the classical brain–lung–thyroid triad for diagnosis. Novel contributions include the delineation of early motor delay as the most typical presenting sign, genotype–phenotype associations that inform motor and respiratory outcomes, and the identification of NRDS as a predictor of later respiratory morbidity. Together, these insights refine prognosis and support a multidisciplinary care model.

Motor delay within the first year was frequent and often preceded other manifestations, occurring independently of global neurodevelopmental delay.<sup>1,29</sup> Although most individuals had some neurodevelopmental difficulty, cognitive and social deficits were less prominent than motor impairment. The combination of early motor delay with hypothyroidism, early pulmonary disease, or hypotonia should heighten suspicion for *NKX2-1*-RD.



**FIG. 4.** Sankey diagram showing the different treatments received for chorea in individuals with *NKX2-1*-related disorders (*NKX2-1*-RD). MFD, methylphenidate; CBZ, carbamazepine; TBZ, tetrabenazine; CZP, clonazepam; VPA, valproate; TRZ, trazodone; HAL, haloperidol; PRG, pregabalin; ARI, aripiprazole; OLZ, olanzapine; LDX, lisdexanfetamine; THP, trihexyphenidyl; NA, Not available; AMT, amantadine; LT4, levothyroxine. [Color figure can be viewed at [wileyonlinelibrary.com](https://onlinelibrary.wiley.com/doi/10.1002/mds.24017)]

Chorea affected the vast majority of patients and usually began in early childhood, then stabilized or improved in most cases – distinct from the progressive course typical of metabolic or neurodegenerative chorea. Complete resolution was not observed here, although Grass et al. reported it in a subset.<sup>29</sup> The observed stabilization or improvement from childhood onward likely reflects natural history in a non-degenerative disorder, although treatment effects (eg, tetrabenazine, levodopa, or thyroid replacement) may modulate the clinical course in individual cases; our registry was not designed to isolate causal treatment effects.

Importantly, improvement in chorea did not parallel other triad features: hypothyroidism persisted and required treatment, and respiratory disease could remain stable or emerge later (eg, interstitial lung disease or asthma). The mechanism of chorea improvement remains unclear; age-related increases in tone within a non-degenerative motor system is a plausible explanation. Although myoclonus was more frequent after the second decade, we did not observe a statistical association with chorea improvement.

Although all the individuals in this cohort had neurological symptoms, this finding should not be interpreted as universal to *NKX2-1*-RD. Referral pathways mainly involved pediatric neurologists, which likely biased

ascertainment toward individuals with neurological presentations and may have missed those with isolated respiratory or thyroid involvement.

NRDS occurred in about one-third of cases and strongly predicted later respiratory symptoms, extending prior observations.<sup>44</sup> A systematic review of 148 individuals showed a respiratory spectrum from NRDS to asthma and interstitial lung disease; nonsense *NKX2-1* SNVs have even been linked to lung cancer.<sup>45</sup> These features are consistent with *NKX2-1* haploinsufficiency affecting surfactant biology, while the endocrine profile underscores its role in thyroid development.<sup>7</sup> Treatments ranged from oxygen and ventilation to transplantation, and long-term outcomes were heterogeneous.<sup>45</sup>

Hypothyroidism was highly prevalent and often diagnosed in infancy, reinforcing the need for early screening. Individuals with mild or compensated hypothyroidism who were not started on levothyroxine showed more severe chorea compared with those receiving treatment or without hypothyroidism, based on retrospective clinical evaluations. This association should be interpreted with caution, because thyroid status was not systematically evaluated before and after treatment initiation, and treatment decisions depended on local clinical judgement. Although thyroid dysfunction in *NKX2-1*-RD is well recognised,<sup>32,33</sup> our cohort revealed underuse of levothyroxine, despite established benefits of early therapy.



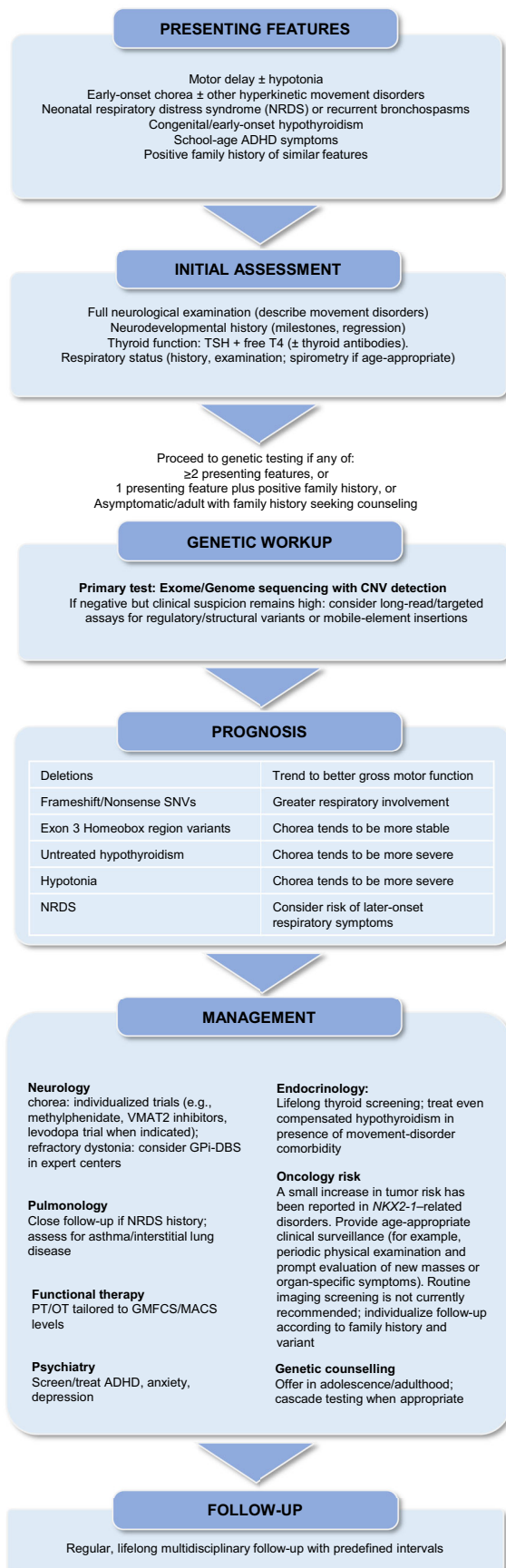


FIG. 5. Legend on next column.

This, together with inconsistent neonatal TSH screening and diagnostic delays, argues for standardized endocrine protocols. We recommend routine thyroid testing in children with neurodevelopmental disorders and lifelong monitoring in all individuals with *NKX2-1*-RD. Treatment of compensated hypothyroidism should be considered carefully given its association with chorea severity, acknowledging the need for confirmation in prospective studies.

Additional motor features – hypotonia, dystonia, and myoclonus – and their interactions (including links between myoclonus, dystonia, and milder neurodevelopmental delay) extend the phenotype beyond chorea. We did not observe consistent dysmorphic features, including in deletion cases, and no malignancies occurred in this relatively young cohort.

Only about half fulfilled the full triad; 40% had dual-system involvement – most often brain–thyroid – and ~9% presented with isolated brain disease. These data reinforce variable expressivity in line with prior reports.<sup>1,29,30,46</sup> The predominance of neurological presentations likely reflects recruitment through neurology and may underrepresent isolated pulmonary or thyroid disease.

Individuals with SNVs were more likely to develop chorea than those with deletions, while deletions – regardless of size or inclusion of neighboring genes – were associated with better gross motor function. Variants in the exon-3 homeobox domain were associated with more stable chorea, whereas non-homeobox variants tended to improve. Frameshift and nonsense variants correlated with increased respiratory involvement, with a near-significant trend for splicing variants. These associations build on earlier work and provide practical guidance for prognosis and follow-up.<sup>43</sup>

The stabilization or improvement of chorea distinguishes *NKX2-1*-RD from progressive disorders such as *DYT-HPCA*, *CHOR/DYT-ADCY5*, and *GNAO1*-related-disease.<sup>47-50</sup> Although there have been descriptions of chorea evolving into myoclonus,<sup>29</sup> our cross-sectional design prevents us from drawing definitive conclusions about longitudinal trajectories. Correlations between

**FIG. 5.** Practical diagnostic and management workflow for *NKX2-1*-related disorders. Algorithm summarizing the clinical approach from initial presentation to genetic work-up, prognosis, and follow-up. Key recommendations include early recognition of motor delay or chorea with neonatal respiratory distress or hypothyroidism, early exome/genome sequencing, and regular multidisciplinary follow-up (neurology, pulmonology, endocrinology, psychiatry, rehabilitation, and genetics). NRDS, neonatal respiratory distress syndrome; ADHD, attention-deficit/hyperactivity disorder; TSH, thyroid-stimulating hormone; T4, free thyroxine; CNV, copy number variant; SNV, single nucleotide variant; VMAT2, vesicular monoamine transporter 2; GPI, globus pallidus internus; DBS, deep brain stimulation; PT, physiotherapy; OT, occupational therapy; GMFCS, Gross Motor Function Classification System; MACS, Manual Ability Classification System. [Color figure can be viewed at [wileyonlinelibrary.com](http://wileyonlinelibrary.com)]

functional motor scales (higher AIMS with worse MACS and GMFCS) and delayed gait with neurodevelopmental severity are novel and suggest that early motor milestones serve as prognostic markers.

We identified Alu retrotransposition in two related individuals and deletions affecting regulatory genes rather than *NKX2-1* itself, expanding the spectrum beyond prior reports and underscoring the value of comprehensive genomic testing.<sup>5</sup>

## Clinical Implications

*NKX2-1*-RD should be considered in neonates with disproportionate NRDS and in infants with isolated motor delay, particularly when hypotonia or early chorea are present, even without overt thyroid disease (Fig. 5). Early genetic testing should interrogate SNVs, deletions, and, when the exome/genome is negative but the phenotype is compelling, regulatory regions and mobile-element insertions. Prognosis should incorporate genotype and sex (eg, better gross motor outcomes with deletions; worse manual ability in males). Differential diagnosis includes *NHLRC2*, which can mimic the respiratory and thyroid phenotype but typically presents with dystonia rather than chorea.<sup>51</sup> The limited and variable response of chorea to medication argues for individualized therapeutic trials, ideally stratified by genotype and sex.

The genotype–phenotype patterns suggest that functional domains of *NKX2-1* differentially influence neurodevelopment, motor control, and respiratory regulation. The relatively milder motor phenotype in deletions versus SNVs may reflect partial preservation of regulatory networks, a hypothesis for future molecular work. Associations between myoclonus, dystonia, and psychiatric comorbidities hint at broader circuit dysfunction, potentially across basal ganglia-thalamo-cortical loops, consistent with contemporary models linking movement disorders and psychiatry.<sup>52</sup>

The cross-sectional design limits causal inference and likely biased recruitment toward neurologically presenting cases; improvement or stabilization was inferred retrospectively from clinical documentation rather than quantified through standardized longitudinal scales, restricting precision in assessing temporal evolution. Although this is the largest cohort to date, subgroup analyses remain underpowered. Management varied across centers, potentially affecting outcomes. The median age at genetic diagnosis suggests a diagnostic delay that may obscure early trajectories. Underrepresentation of pulmonology expertise and interstitial lung disease cases limits generalizability to those subgroups. Severity and longitudinal changes in chorea and myoclonus were derived from retrospective expert clinical documentation and AIMS ratings when available. Because

AIMS lacks validation in *NKX2-1*-RD and standardized serial scales were inconsistently applied at fixed intervals, measurement variability is possible, preventing time-to-event analyses. Additionally, because the registry relied on clinician notes rather than prospective standardized assessments, the longitudinal evolution of chorea may be subject to documentation bias.

Prospective, longitudinal cohorts with standardized neurological, pulmonary, and endocrine assessments are needed to validate predictors (eg, NRDS for later pulmonary disease) and to define natural history. Mechanistic studies contrasting homeobox versus non-homeobox SNVs and assessing regulatory disruptions may reveal therapeutic targets. Genotype- and sex-stratified treatment studies, together with routine psychiatric screening, should inform evidence-based care.

In summary, *NKX2-1*-RD is clinically heterogeneous, but chorea, NRDS, and hypothyroidism remain defining features. Motor delay is a frequent early sign that may presage chorea. By mapping genotype-specific outcomes and early clinical predictors, this study offers a framework for genotype-informed diagnosis, counseling and management, reinforcing the need for coordinated care across neurology, pulmonology, endocrinology, and psychiatry. ■

**Author Roles:** (1) Research Project: A. Conceptualization, B. Methodology, C. Data Curation, D. Investigation, E. Data Analysis, F. Visualization; (2) Statistical Analysis: A. Design, B. Execution, C. Review and Critique; (3) Manuscript Preparation: A. Writing of the First Draft, B. Review and Editing; (4) Other: A. Supervision.

L.N.F.: 1B, 1C, 1D, 1E, 1F, 3A, 3B.  
S.B.M., A.V.-V., E.R.S.: 1C, 1D, 1E, 1F, 3B.  
C.R., B.V., A.S.-V., C.V., R.B., S.N., D.N.d.B., D.C.-A., L.S., C.M.d.G., G.G., F.M., C.C., R.P., E.R., D.D.: 1C, 1D, 3B.  
L.B., A.I., V.Q., A.S.J., G.B., A.D.-G., A.A., P.D.d.S.M., D.G.-N.N., M.K., M.J.M., L.M., K.O., M.P., P.V., M.V.-O., N.S., M.C., J.C.M., K.B., D.E.-F., M.K., J.N.C.: 1C, 1D, 3B.  
J.D.O.-E.: 1A, 1B, 1C, 1D, 1F, 3B, 4A.

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## 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.

## References

1. Thorwarth A, Schnittert-Hübener S, Schrupf P, et al. Comprehensive genotyping and clinical characterisation reveal 27 novel

- NKX2-1 mutations and expand the phenotypic spectrum. *J Med Genet* 2014;51(6):375–387. <https://doi.org/10.1136/jmedgenet-2013-102248>
2. Nattes E, Lejeune S, Carsin A, et al. Heterogeneity of lung disease associated with NK2 homeobox 1 mutations. *Respir Med* 2017;129:16–23. <https://doi.org/10.1016/j.rmed.2017.05.014>
3. Yang L, Lin M, Ruan WJ, et al. Nkx2-1: a novel tumor biomarker of lung cancer. *J Zhejiang Univ Sci B* 2012;13(11):855–866. <https://doi.org/10.1631/jzus.B1100382>
4. Möller K, Gulzar T, Lennartz M, et al. TTF-1 is a highly sensitive but not fully specific marker for pulmonary and thyroid cancer: a tissue microarray study evaluating more than 17,000 tumors from 152 different tumor entities. *Virchows Arch* 2024;485(5):815–828. <https://doi.org/10.1007/s00428-024-03926-1>
5. Magrinelli F, Rocca C, Simone R, et al. Detection and characterization of a de novo alu retrotransposition event causing NKX2-1-related disorder. *Mov Disord* 2023;38(2):347–353. <https://doi.org/10.1002/mds.29280>
6. Liao J, Coffman KA, Locker J, et al. Deletion of conserved non-coding sequences downstream from NKX2-1: a novel disease-causing mechanism for benign hereditary chorea. *Mol Genet Genomic Med* 2021;9(4):1–11. <https://doi.org/10.1002/mgg3.1647>
7. Kimura S, Hara Y, Pineau T, et al. The T/ebp null mouse: thyroid-specific enhancer-binding protein is essential for the organogenesis of the thyroid, lung, ventral forebrain, and pituitary. *Genes Dev* 1996;10(1):60–69. <https://doi.org/10.1101/gad.10.1.60>
8. Minoo P, Hamdan H, Bu D, Warburton D, Stepanik P, Delemos R. TTF-1 regulates lung epithelial morphogenesis. *Dev Biol* 1995;172(2):694–698. <https://doi.org/10.1006/dbio.1995.8080>
9. Little DR, Lynch AM, Yan Y, Akiyama H, Kimura S, Chen J. Differential chromatin binding of the lung lineage transcription factor NKX2-1 resolves opposing murine alveolar cell fates in vivo. *Nat Commun* 2021;12(1):1–18. <https://doi.org/10.1038/s41467-021-22817-6>
10. Bruno MD, Bohinski RJ, Huelsman KM, Whitsett JA, Korfhagen TR. Lung cell-specific expression of the murine surfactant protein A (SP-A) gene is mediated by interactions between the SP-A promoter and thyroid transcription factor-1. *J Biol Chem* 1995;270(12):6531–6536. <https://doi.org/10.1074/jbc.270.12.6531>
11. Kelly SE, Bachurski CJ, Burhans MS, Glasser SW. Transcription of the lung-specific surfactant protein C gene is mediated by thyroid transcription factor 1. *J Biol Chem* 1996;271(12):6881–6888. <https://doi.org/10.1074/jbc.271.12.6881>
12. Ikeda K, Clark JC, Shaw-White JR, Stahlman MT, Boutell CJ, Whitsett JA. Gene structure and expression of human thyroid transcription factor-1 in respiratory epithelial cells. *J Biol Chem* 1995;270(14):8108–8114. <https://doi.org/10.1074/jbc.270.14.8108>
13. Civitareale D, Lonigro R, Sinclair AJ, Di Lauro R. A thyroid-specific nuclear protein essential for tissue-specific expression of the thyroglobulin promoter. *EMBO J* 1989;8(9):2537–2542. <https://doi.org/10.1002/f.1460-2075.1989.tb08391.x>
14. Tübing J, Bohnenpoll J, Spiegler J, et al. Methylphenidate can improve chorea in NKX2.1 and ADCY5 mutation-positive patients – a report of two children. *Mov Disord Clin Pract* 2018;5(3):343–345. <https://doi.org/10.1002/mdc3.12608>
15. Koht J, Løstegaard SO, Wedding I, Vidailhet M, Louha M, Tallaksen CME. Benign hereditary chorea, not only chorea: a family case presentation. *Cerebellum Ataxias* 2016;3(1):1–7. <https://doi.org/10.1186/s40673-016-0041-7>
16. Ferrara JM, Adam OR, Kirwin SM, et al. Brain-lung-thyroid disease: clinical features of a kindred with a novel thyroid transcription factor 1 mutation. *J Child Neurol* 2012;27(1):68–73. <https://doi.org/10.1177/0883073811413584>
17. Parnes M, Bashir H, Jankovic J. Is benign hereditary chorea really benign? Brain-lung-thyroid syndrome caused by NKX2-1 mutations. *Mov Disord Clin Pract* 2019;6(1):34–39. <https://doi.org/10.1002/mdc3.12690>
18. Sempere AP, Aparicio S, Mola S, Pérez-Tur J. Benign hereditary chorea: clinical features and long-term follow-up in a Spanish family. *Parkinsonism Relat Disord* 2013;19(3):394–396. <https://doi.org/10.1016/j.parkreldis.2012.08.006>
19. Konishi T, Kono S, Fujimoto M, et al. Benign hereditary chorea: dopaminergic brain imaging in patients with a novel intronic NKX2.1 gene mutation. *J Neurol* 2013;260(1):207–213. <https://doi.org/10.1007/s00415-012-6618-z>
20. Santos-Silva R, Rosário M, Grangeira A, et al. Genetic analyses in a cohort of Portuguese pediatric patients with congenital hypothyroidism. *J Pediatr Endocrinol Metab* 2019;32(11):1265–1273. <https://doi.org/10.1515/jpem-2019-0047>
21. Williamson S, Kirkpatrick M, Greene S, Goudie D. A novel mutation of NKX2-1 affecting 2 generations with hypothyroidism and choreoathetosis: part of the spectrum of brain-thyroid-lung syndrome. *J Child Neurol* 2014;29(5):666–669. <https://doi.org/10.1177/0883073813518243>
22. Balicza P, Grosz Z, Molnár V, et al. NKX2-1 new mutation associated with myoclonus, dystonia, and pituitary involvement. *Front Genet* 2018;9:1–5. <https://doi.org/10.3389/fgene.2018.00335>
23. Asmus F, Devlin A, Munz M, Zimprich A, Gasser T, Chinnery PF. Clinical differentiation of genetically proven benign hereditary chorea and myoclonus-dystonia. *Mov Disord* 2007;22(14):2104–2109. <https://doi.org/10.1002/mds.21692>
24. Veneziano L, Parkinson MH, Mantuano E, Frontali M, Bhatia KP, Giunti P. A novel de novo mutation of the TITF1/NKX2-1 gene causing ataxia, benign hereditary chorea, hypothyroidism and a pituitary mass in a UK family and review of the literature. *Cerebellum* 2014;13(5):588–595. <https://doi.org/10.1007/s12311-014-0570-7>
25. Devos D, Vuillaume I, de Believre A, et al. New syndromic form of benign hereditary chorea is associated with a deletion of TITF-1 and PAX-9 contiguous genes. *Mov Disord* 2006;21(12):2237–2240. <https://doi.org/10.1002/mds.21135>
26. Salvatore E, di Maio L, Filla A, et al. Benign hereditary chorea: clinical and neuroimaging features in an Italian family. *Mov Disord* 2010;25(10):1491–1495. <https://doi.org/10.1002/mds.23065>
27. Rosati A, Berti B, Melani F, Cellini E, Procopio E, Guerrini R. Recurrent drop attacks in early childhood as presenting symptom of benign hereditary chorea caused by TITF1 gene mutations. *Dev Med Child Neurol* 2015;57(8):777–779. <https://doi.org/10.1111/dmcn.12644>
28. de Gusmao CM, Kok F, Casella EB, Waugh JL. Benign hereditary chorea related to NKX2-1 with ataxia and dystonia. *Neurol Genet* 2016;2(1):1–2. <https://doi.org/10.1212/NXG.0000000000000040>
29. Gras D, Jonard L, Roze E, et al. Benign hereditary chorea: phenotype, prognosis, therapeutic outcome and long term follow-up in a large series with new mutations in the TITF1/NKX2-1 gene. *J Neurol Neurosurg Psychiatry* 2012;83(10):956–962. <https://doi.org/10.1136/jnnp-2012-302505>
30. Peall KJ, Lumsden D, Kneen R, et al. Benign hereditary chorea related to NKX2.1: expansion of the genotypic and phenotypic spectrum. *Dev Med Child Neurol* 2014;56(7):642–648. <https://doi.org/10.1111/dmcn.12323>
31. Nou-Fontanet L, Martín-Gómez C, Isabel-Gómez R, et al. Systematic review of drug therapy for chorea in NKX2-1-related disorders: efficacy and safety evidence from case studies and series. *Eur J Neurol* 2023;30(12):3928–3948. <https://doi.org/10.1111/ene.16038>
32. Carmona-Hidalgo B, Martín-Gómez C, Herrera-Ramos E, et al. Systematic review of thyroid function in NKX2-1-related disorders: screening and diagnosis. *PLoS One* 2024;19(7 July):1–17. <https://doi.org/10.1371/journal.pone.0303880>
33. Carmona-Hidalgo B, Herrera-Ramos E, Rodríguez-López R, et al. Systematic review of thyroid function in NKX2-1-related disorders: treatment and follow-up. *PLoS One* 2024;19(10):e0309064. <https://doi.org/10.1371/journal.pone.0309064>
34. Invernizzi F, Zorzi G, Legati A, et al. Benign hereditary chorea and deletions outside NKX2-1: What's the role of MBIP? *Eur J Med Genet* 2018;61(10):581–584. <https://doi.org/10.1016/j.ejmg.2018.03.011>
35. Santagati F, Abe K, Schmidt V, et al. Identification of cis-regulatory elements in the mouse Pax9/Nkx2-9 genomic region: implication for evolutionary conserved synteny. *Genetics* 2003;165(1):235–242. <https://doi.org/10.1093/genetics/165.1.235>

36. Harris PA, Taylor R, Thielke R, Payne J, Gonzalez N, Conde JG. Research electronic data capture (REDCap)-a metadata-driven methodology and workflow process for providing translational research informatics support. *J Biomed Inform* 2009;42(2):377–381. <https://doi.org/10.1016/j.jbi.2008.08.010>
37. Harris PA, Taylor R, Thielke R, et al. The REDCap consortium: building an International Community of Software Platform Partners. *J Biomed Inform* 2009;42(2):377–381. <https://doi.org/10.1016/j.jbi.2019.103208>
38. Skwara J, Nowicki M, Sharif L, et al. Differential diagnosis of Huntington's disease – neurological aspects of NKX2-1-related disorders. *J Neural Transm* 2024;131(9):1013–1024. <https://doi.org/10.1007/s00702-024-02800-3>
39. Graziola F, Garone G, Grasso M, Schirizzi T, Capuano A. Working memory, attention and planning abilities in NKX2.1-related chorea. *Parkinsonism Relat Disord* 2021;88:24–27. <https://doi.org/10.1016/j.parkreldis.2021.05.021>
40. Fons C, Rizzu P, Garcia-Cazorla A, et al. TITF-1 gene mutation in a case of sporadic non-progressive chorea. Response to levodopa treatment. *Brain Dev* 2012;34(3):255–257. <https://doi.org/10.1016/j.braindev.2011.04.007>
41. Puusepp S, Reinson K, Pajusalu S, et al. Effectiveness of whole exome sequencing in unsolved patients with a clinical suspicion of a mitochondrial disorder in Estonia. *Mol Genet Metab Rep* 2018;15: 80–89. <https://doi.org/10.1016/j.ymgmr.2018.03.004>
42. Villafuerte B, Natera-de-Benito D, González A, et al. The brain-lung-thyroid syndrome (BLTS): a novel deletion in chromosome 14q13.2-q21.1 expands the phenotype to humoral immunodeficiency. *Eur J Med Genet* 2018;61(7):393–398. <https://doi.org/10.1016/j.ejmg.2018.02.007>
43. Villafuerte B, Carrasco-López C, Herranz A, et al. A novel missense variant in the NKX2-1 homeodomain prevents transcriptional rescue by TAZ. *Thyroid* 2024;34(7):942–948. <https://doi.org/10.1089/thy.2023.0593>
44. Hamvas A, Deterding RR, Wert SE, et al. Heterogeneous pulmonary phenotypes associated with mutations in the thyroid transcription factor gene NKX2-1. *Chest* 2013;144(3):794–804. <https://doi.org/10.1378/chest.12-2502>
45. Michel K, Ruiz-Ramos A, Nou-Fontanet L, et al. Respiratory and other organ manifestations in NKX2-1-related disorders: a systematic review. *Front Med* 2025;12:1–13. <https://doi.org/10.3389/fmed.2025.1507513>
46. Carré A, Szinnai G, Castanet M, et al. Five new TTF1/NKX2.1 mutations in brain-lung-thyroid syndrome: rescue by PAX8 synergism in one case. *Hum Mol Genet* 2009;18(12):2266–2276. <https://doi.org/10.1093/hmg/ddp162>
47. Breedveld GJ, van Dongen JWF, Danesino C, et al. Mutations in TITF-1 are associated with benign hereditary chorea. *Hum Mol Genet* 2002;11(8):971–979. <https://doi.org/10.1093/hmg/11.8.971>
48. Lange LM, Gonzalez-Latapi P, Rajalingam R, et al. Nomenclature of genetic movement disorders: recommendations of the International Parkinson and Movement Disorder Society Task Force – an update. *Mov Disord* 2022;37(5):905–935. <https://doi.org/10.1002/mds.28982>
49. Pérez-Dueñas B, Gorman K, Marcé-Grau A, et al. The genetic landscape of complex childhood-onset hyperkinetic movement disorders. *Mov Disord* 2022;37(11):2197–2209. <https://doi.org/10.1002/mds.29182>
50. Domínguez Carral J, Reinhard C, Ebrahimi-Fakhari D, et al. Dyskinetic crisis in GNAO1-related disorders: clinical perspectives and management strategies. *Front Neurol* 2024;15:1–11. <https://doi.org/10.3389/fneur.2024.1403815>
51. Tallgren A, Kager L, Grady GO, et al. Novel patients with NHLRC2 variants expand the phenotypic spectrum of FINCA disease. *Front Neurosci* 2023;15:1123327. <https://doi.org/10.3389/fnins.2023.1123327>
52. Malt EA, Juhasz K, Malt UF, Naumann T. A role for the transcription factor Nk2 homeobox 1 in schizophrenia: convergent evidence from animal and human studies. *Front Behav Neurosci* 2016;10:59. <https://doi.org/10.3389/fnbeh.2016.00059>

## Supporting Data

Additional Supporting Information may be found in the online version of this article at the publisher's web-site.