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The Genetic Basis of Incomitant Strabismus: Consolidation of the Current Knowledge of the Genetic Foundations of Disease:

The Genetic Basis of Incomitant Strabismus

Carolyn P. Graeber, MD1,3, **David G. Hunter, MD, PhD**1,3, and **Elizabeth C. Engle, MD**1,2,3,4 Department of Ophthalmology, Boston Children's Hospital, 300 Longwood Ave, Boston, MA Department of Neurology, Boston Children's Hospital, 300 Longwood Ave, Boston, MA Department of Ophthalmology, Harvard Medical School, 243 Charles St, Boston, MA Department of Neurology, Harvard Medical School, 243 Charles St, Boston, MA

Abstract

In recent years, our understanding of the genetic foundations of incomitant strabismus has grown significantly. Much new understanding has been gleaned since the concept of congenital cranial dysinnervation disorders (CCDDs) was introduced in 2002, and the genetic basis of CCDDs continues to be elucidated. In this review, we aim to provide an update of the genetic and clinical presentation of these disorders. Disorders reviewed include Duane syndrome (DS), *HOXA1* and *HOXB1* syndromes, Moebius syndrome, congenital fibrosis of the extraocular muscles (CFEOM), and horizontal gaze palsy with progressive scoliosis (HGPPS).

Keywords

congenital cranial dysinnervation disorders; congenital fibrosis of the extraocular muscles; Duane syndrome; Moebius syndrome; horizontal gaze palsy with progressive scoliosis; *HOX* mutations

Introduction

Congenital cranial dysinnervation disorders (CCDDs) are a group of disorders that result from the abnormal development of cranial motor nuclei and their respective cranial nerves. Because these cranial nerves fail to appropriately innervate their intended target muscles, these muscles often have fibrotic changes, which is now understood to be secondary to either absent or aberrant innervation from other nearby cranial nerves.

Our understanding of the CCDDs has evolved over time. The path to understanding the history of this is best exemplified by congenital fibrosis of the extraocular muscles (CFEOM). Long recognized as a source of ocular pathology, elements of this disorder were first described as early as the late 1800s, but it was not until 1956 that Laughlin first coined the term CFEOM.¹ The basis of pathology was initially thought to be the muscles themselves, which showed fibrosis histologically and restrictive strabismus clinically.

Corresponding Author: Elizabeth C. Engle, MD, 300 Longwood Ave, Boston, MA, elizabeth.engle@childrens.harvard.edu.

Through the 1980s and 90s, however, mounting evidence showed that the underlying pathology was the agenesis or hypoplasia of the oculomotor nerve and that the disorder was in fact absent or aberrant innervation of these muscles that caused secondary fibrotic and restrictive changes. Other disorders were also shown to have similar underlying pathogenesis, including Duane syndrome (DS) and Moebius syndrome. This new understanding prompted the reclassification of these disorders under the umbrella term "CCDDs."

In this review, we discuss the current understanding of the genetic underpinnings of CCDDs. Table 1 summarizes the disorders, their clinical and radiographic findings, and the genes that have been associated with each of them.

Methods

A literature search of PubMed (covering years 1950-January, 2013) for the following terms was conducted: "congenital cranial dysinnervation disorders," "Duane syndrome," "*CHN1* and Duane syndrome," "Duane radial ray syndrome," "*CHN1* and Duane syndrome," "Goldenhar and Duane syndrome," "*SALL1* and Duane syndrome," "8q12 duplication and Duane syndrome," "8q13 and Duane syndrome," "Wildervanck and Duane syndrome," "*HOXA1* and Bosley-Salih-Alorainy syndrome," "*HOXA1* and Athabascan Brain Dysgenesis syndrome," "Moebius syndrome," "*HOXB1* and Moebius syndrome," "congenital fibrosis of the extraocular muscles," "congenital fibrosis of the extraocular muscles and *KIF21A*," "CFEOM and *PHOX2A*," "CFEOM and *TUBB3*," "horizontal gaze palsy with progressive scoliosis," and "horizontal gaze palsy with progressive scoliosis and *ROBO3*." Reference lists within pertinent articles were reviewed. Only English language papers with relevance to ocular findings were reviewed.

Isolated (Non-Syndromic) Duane Syndrome

The most common CCDD, Duane syndrome (DS), has been reported to account for 1–4% of strabismus cases.^{2,3} DS is characterized by congenital absence or hypoplasia of the abducens nucleus and subsequent aberrant innervation of the lateral rectus muscle by branches of the oculomotor nerve.2,3 Clinically, this syndrome manifests as decreased horizontal movement of the affected eye with narrowing of the palpebral fissure and globe retraction with attempted adduction.⁴

Three types of DS have been described. In type 1 DS, the affected eye has limited abduction but preserved adduction, frequently resulting in esotropia in primary gaze. In type 2, the affected eye has limited adduction but preserved abduction, often causing exotropia in primary gaze. In type 3, the affected eye has limitation of both abduction and adduction. In many patients, these types are not distinct once the eye movements have been carefully scrutinized. The primary gaze deviation can be esotropia, exotropia, or orthophoria, depending on the balance between the deficits of the horizontal muscles.

Most individuals with Duane syndrome are the only affected member of their family (simplex, or sporadic, cases), but hereditary forms account for $5-10\%$ of cases.⁵ Several investigators have described bilateral DS with an autosomal dominant inheritance pattern.^{6,7}

Recent mapping of the phenotype of families with DS inherited as a dominant trait identified the DURS2 locus on chromosome 2 and, subsequently, heterozygous mutations in *CHN1*. 5,8–12 Current evidence suggests that gain of function mutations in *CHN1* hyperactivate α2-chimerin and cause disruption in the growth or guidance of cranial axons destined to innervate extraocular muscles during development.^{11,12}

Phenotypic variability has been recognized among individuals harboring heterozygous *CHN1* mutations.7,11,12 Many affected individuals have bilateral DS with either manifestations of either type 1 or type 3 DS. Some have one type 1 in one eye and type 3 in the other, but none have type 2.⁹ In addition, a subset also has vertical deviations associated with DS, $9,12-14$ and some have vertical deviations in the absence of DS.¹² Given these features, it is reasonable for patients with bilateral DS and associated vertical motility anomalies^{12,14} and patients with familial vertical deviations¹² to be screened for *CHN1* mutations. Individuals harboring the familial mutation do not always have clinical evidence of disease, indicating that while phenotypic penetrance is very high, it is not complete.^{14,15}

Demer and colleagues performed magnetic resonance imaging (MRI) on 8 family members from two autosomal dominant DS pedigrees who harbored *CHN1* mutations and compared the findings with those of 11 control patients who had comitant strabismus. The DS patients showed a variable endophenotype, with most participants showing markedly abnormal lateral rectus muscles and some showing abnormalities in other extraocular muscles as well. The only muscles that were not affected were those supplied by the inferior division of the oculomotor nerve: the inferior rectus, medial rectus, and inferior oblique muscles. This finding suggests that the abnormality in DS may not be limited to just the abducens nerve but may also involve the superior branch of the oculomotor nerve. In addition, there was evidence of superior oblique hypoplasia in half of the individuals tested, suggesting that the trochlear nerve may also be affected in some individuals.¹³

It is important to note that *CHN1* has not been identified as a cause of simplex $DS¹⁵$ nor has it been implicated in Brown syndrome, congenital disorders of the oblique muscles, or vertical retraction syndrome.¹⁶ *CHN1* hyperactivation has, however, been associated with deficits in supraduction in the absence of DS.¹² Abnormalities in chromosomes 10 and 22 have also been implicated in the pathogenesis of $DS¹⁷$ Therefore, while much information on the genetic basis of DS has been gleaned in recent years, much more remains to be determined.

Syndromic Duane Syndrome

While the majority of DS occurs in isolation, DS is associated with characteristic systemic findings in about 30% of cases. In this section, we describe the clinical features and genetic analysis to date of these syndromes.

A spectrum of overlapping disorders that include DS with radial limb abnormalities, facial asymmetry, hearing deficits, ear abnormalities, anal stenosis, and cardiac and renal abnormalities have been associated with mutation in *SALL4*. 19,20 Individuals with *SALL4* mutations have also been found in radial limb anomalies without DS.18 Disorders on this spectrum include Duane radial ray (Okihiro) syndrome, Holt-Oram syndrome, and acro-

renal-ocular syndrome, all of which have been shown to have links to *SALL4* mutations.19 A *SALL4* mutation has also been identified in a man previously thought to have thalidomide embryopathy who then had a daughter with similar limb deformities.¹⁹ MRI analysis of a family with a heterozygous *SALL4* mutation resulting in Duane radial ray syndrome revealed hypoplastic to absent abducens nerves with normal intracranial oculomotor and optic nerves. In some subjects, a branch of the oculomotor nerve was shown to be proximal to the lateral rectus muscle, implying that it may partially innervate this muscle.¹³ *SALL4* mutations have not been identified in patients with simplex $DS^{21,22}$ Table 2a summarizes the clinical differences among *SALL4* disorders.

Townes-Brocks syndrome is an autosomal dominant condition that is associated with mutations in *SALL1*. ²⁰ This syndrome is characterized by ear, limb, anal, and renal anomalies but has rarely been associated with DS.²¹

Abnormalities on chromosome 8 have been observed to co-segregate with DS. A reciprocal balanced translocation in chromosome 8q13 implicated *CPAH* as a possible etiology of simplex DS.²² Three other patients with disruption of this gene locus have exhibited DS with various other systemic associations.^{23–25} Chromosome 8q12 duplications have also been reported to produce a combination of DS with sensorineural deafness, cardiac defects, hypotonia, and developmental delay.26 Duplications in this chromosome locus can also produce developmental delay and particular facial features, including full cheeks, a specific lip shape, and horizontal and flared evebrows, with or without $DS^{27,28}$ The duplicated region of 8q12 includes *CHD7*, implying that abnormal dosage of the transcribed protein may be a factor.^{26,27}

Wildervanck syndrome (also known as cervico-oculo-acoustic syndrome) is characterized by Duane syndrome, Klippel-Feil deformity of the spine (congenitally fused cervical vertebrae), and hearing loss. This syndrome has a female predominance, suggesting a possible defect on the X chromosome that is lethal in hemizygous males.²⁹ Genetic evaluation of a simplex case in an affected male revealed a microdeletion on chromosome X involving only a single gene: Fibroblast Growth Factor Homologous Factor 13 (*FGFHF13*).³⁰

Goldenhar syndrome, 31 which occurs in as many as 3% of DS patients, 32 has been associated with deletions in chromosome $22q11.2$ ^{33,34} While some DS has been linked to chromosome 22, the exact locus responsible for the association of these two disorders is, as yet, unknown. DS has been linked to numerous other defined conditions in isolated case reports (e.g., Marfan syndrome), 33 but to date, these do not have an identified genetic basis.

HOXA1 Syndromes

Recessive, homozygous *HOXA1* mutations have been reported to cause horizontal gaze palsy, facial weakness, and sensorineural hearing loss sometimes associated with developmental delay and hypoventilation.³⁴ Initially, two syndromes, Bosley-Salih-Alorainy Syndrome (BSAS) and Athabascan Brain Dysgenesis syndrome (ABDS), were described. BSAS was described in the Middle Eastern population and included bilateral type 3 DS, sensorineural hearing loss, malformation of the cerebral vasculature, and autism in select

patients.34 MRI findings showed grossly normal orbits and extraocular muscles. The internal carotid artery, however, was hypoplastic or absent in all individuals, and a few patients had duplication of the vertebral artery. The abducens nerve appeared to be absent bilaterally in one patient.35 The second variant, ABDS, was reported in the Native American population. These individuals had horizontal gaze restriction, facial weakness, hearing impairment, cerebral vasculature malformation, cardiac malformations, central hypoventilation, and intellectual disabilities.34,36

Despite some initial differences noted between BSAS and ABDS, overlapping features between the two populations were noted at the outset.³⁴ Subsequently, members of a BSAS family were reported to also have cardiac anomalies (a feature more in line with ABDS), and members of the ABDS family had mild cognitive changes more consistent with BSAS. The study also found that not all patients harboring a homozygous *HOXA1* mutation had horizontal gaze limitation or deafness.³⁷ These findings further blurred the distinction between the two entities and widen the *HOXA1* human phenotype.

Homozygous *HOXA1* mutations result in phenotypes that overlap with type 3 DS and Moebius syndrome (see section on Moebius syndrome below). Evaluation of *HOXA1* in DS and Moebius patients, however, did not reveal any mutations, indicating that *HOXA1* mutations are not a major cause of simplex DS^{38} or Moebius syndrome.³⁹ Table 2b summarizes the clinical differences between ABDS and BSAS.

Moebius Syndrome

Moebius syndrome is a nonprogressive, congenital facial paralysis with limited abduction of the eyes; it can be associated with hearing impairment and other cranial nerve dysfunction, as well as other developmental abnormalities.40 The syndrome occurs in simplex cases and is more likely to occur in children exposed to misoprostol in utero.⁴¹ While some mutations have been identified in patients with atypical forms of Moebius syndrome, no mutations have been observed in patients with typical findings. The genetic basis of this syndrome, if one exists, remains elusive.

Carta and colleagues⁴⁵ evaluated a large cohort of patients with this syndrome to identify their clinical characteristics. Three different patterns of eye movements were identified. The most common motility pattern, identified in half of the patients, was a large esotropia with almost complete abduction deficit but relatively conserved adduction. These patients utilized cross-fixation to compensate for their duction deficits. The next most common ocular motility pattern was orthotropia in primary gaze with deficits of abduction and adduction, requiring large head movements to look to either side. Vertical ductions were preserved. Finally, the smallest number of patients (less than 10%) exhibited a large angle exotropia with absence of convergence. These patients also had limited vertical gaze accompanied by torticollis.⁴²

HOXB1

Identified in a conservative German American population, a single homozygous *HOXB1* mutation has been associated with comitant strabismus, bilateral facial palsy, and hearing

impairment in two families. While affected individuals had facial palsy and esotropia, they did not meet criteria for Moebius syndrome, as they had full abduction of both eyes. The mutation is predicted to result in loss of *HOXB1* function.⁴³

Congenital Fibrosis of the Extraocular Muscles

Congenital fibrosis of the extraocular muscles (CFEOM) is a congenital, non-progressive restrictive ophthalmoplegia and ptosis that has been shown to have a neurogenic basis. Three phenotypes have been described: CFEOM1, 2, and 3.

CFEOM1, the most common of the three phenotypes, is defined by the combination of nonprogressive, bilateral infraduction of the eyes in resting position, limited vertical eye movements with the inability to elevate either eye over the horizontal midline, and bilateral ptosis with a compensatory "chin up" head posture (Figure 1). Many individuals have been noted to have "A pattern" strabismus.⁴⁴ Pupil size and response are normal. Autopsy and radiographic studies have shown hypoplasia or agenesis of the oculomotor nerve, in particular its superior division.^{46,47} These studies support the hypothesis that the primary pathology of CFEOM is dysinnervation, with secondary fibrosis of the target muscles.⁴⁵

Inheritance of CFEOM1 is autosomal dominant. Mutations in *KIF21A* on chromosome 12, which encodes a kinesin motor protein, have been identified in most individuals with CFEOM1.46,47 Rare CFEOM1 patients harbor mutations in *TUBB3*, ⁴⁸ and other genes not yet identified are likely to be responsible for CFEOM1 in a minority of individuals.49 In addition, in rare individuals, mutations in *KIF21A* result in clinical findings most consistent with CFEOM3 (see description below).⁵⁰

Individuals with CFEOM2 have profound ptosis, exotropia, and small, poorly reactive pupils.51 The condition was found to result from homozygous mutations in *PHOX2A*, 52 which encodes a homeodomain transcription factor. To date, there has been no evidence of *PHOX2A* mutations in patients with CFEOM1 or $3,53,54$ indicating that this is a genetically distinct entity.

Clinical evaluation of patients with *PHOX2A* mutations found ptosis often with compensatory chin-up position, pupils that were unreactive to light but that retained response to pharmacologic agents, and large angle exotropia with compensatory head turn away from the fixing eye.55 Convergence and abduction were also almost absent. Neuroimaging revealed large lateral rectus muscles consistent with the exotropic eye position, with all other rectus muscles comparatively small. These studies suggest that the clinical findings in CFEOM2 are the result of isolated but complete dysinnervation of targets of the oculomotor and trochlear nerves, with preservation of all other cranial nerves.

CFEOM3 is a third phenotype that has variable clinical findings that include varying degrees of ptosis and ophthalmoplegia, ranging from mild to complete. In the severest form, these patients have profound ptosis and abducted and infraducted position of the globes with severe restriction of motility (Figure 2). In mildly affected individuals, the only deficit is decreased supraduction with the globes in a normal position in primary gaze. Findings can

be unilateral or bilateral. Some of these individuals resemble CFEOM1 or CFEOM2 phenotypes. Familial transmission is autosomal dominant with variable penetrance.

CFEOM3 can result from heterozygous missense mutations in *TUBB3*, which encodes a βtubulin isotype that is a component of neuronal microtubules.48 While some *TUBB3* mutations result in isolated CFEOM3, other specific mutations cause dysfunction of additional cranial and spinal nerves, and social and intellectual disabilities accompanied by maldevelopment of the corpus callosum and morphologic changes in the basal ganglia.⁴⁸

One specific *TUBB3* mutation results in an E410K amino acid substitution and results in what is now referred to as the *TUBB3* E410K syndrome.³⁹ Clinical overlap exists between atypical Moebius syndrome and the *TUBB3* E410K syndrome. Seven of the eight subjects found to have this mutation previously had a diagnosis of Moebius syndrome until genetic evidence showed *de novo* heterozygous missense *TUBB3* mutations.39 These individuals clinically had ptosis and limited upgaze, more consistent with CFEOM3 than with typical Moebius syndrome.

MRI findings in patients with *TUBB3* mutations reveal structural correlations with the clinical phenotypes. In individuals with severe phenotypes that clinically resemble CFEOM1, there is hypoplasia of the superior rectus, medial rectus, and levator palpebrae superioris muscles. In some cases, the inferior oblique muscle is also hypoplastic. Aberrent branches of the oculomotor nerve suggest misinnervation of the lateral rectus muscle. Individuals with milder phenotypes also had fewer abnormalities on MRI imaging, with relatively less hypoplasia of the subarachnoid portion of the oculomotor nerve and its targeted extraocular muscles.⁵⁶

Recently, a homozygous missense *TUBB2B* mutation was found to be associated with CFEOM and polymicrogyria. To date this is the only mutation in *TUBB2B* that has been associated with a CFEOM.57 In addition, rare individuals meeting CFEOM3 criteria harbor *KIF21A* mutations.53,58,59

While the above genes account for the majority of families with CFEOM, additional mutations causing the disorder are likely yet to be identified and characterized. One mutation-negative pedigree exhibits the phenotype of CFEOM3 and has a translocation that implicates a locus on chromosome $13q12$ (FEOM4) as the source of pathogenesis.⁶⁰ Additional autosomal recessive forms of CFEOM also appear to exist. A consanguineous Turkish family with autosomal recessive transmission of a unilateral, non-progressive ophthalmoplegia and hand abnormalities mapped to chromosome 21 ;⁶¹ however, forced ductions were reportedly normal in this family, suggesting an atypical presentation for a CFEOM. In one Lebanese family, affected members had unilateral ptosis and restrictive strabismus but were found to have no identifiable linkage to the CFEOM1, -2, or -3 locus and had no mutation in *KIF21A* or *PHOX2A* genes.⁶² The full spectrum of mutations that can give rise to CFEOM phenotype, therefore, remains to be elucidated.

Horizontal Gaze Palsy with Progressive Scoliosis

First described in 1975, horizontal gaze palsy with progressive scoliosis (HGPPS) is a rare autosomal recessive disorder that features lateral gaze limitation and scoliosis (Figure 3) as key clinical findings.63 It was not until 2004, however, that the pathogenesis was linked to mutations in *ROBO3* in consanguineous families with autosomal recessive inheritance pattern of the disease.⁶⁴ This gene is necessary for hindbrain axons to appropriately cross the midline. Corresponding defects in MRI imaging of the hindbrain were identified in affected individuals, including a characteristic anterior flattening and midline cleft now referred to as a butterfly configuration on axial cuts (Figure 4).⁶⁴ HGPPS can also result from compound heterozygous mutations in non-consanguineous families.65 Presenting signs in these patients were plagiocephaly and torticollis, both of which developed before scoliosis. Therefore, infants and children presenting with plagiocephaly or torticollis in addition to lateral gaze palsy would benefit from genetic testing for *ROBO3* mutations.⁶⁵ Because of the ophthalmologic and molecular cues that can help diagnose HGPPS, patients can be diagnosed from an early age and screened for the onset of scoliosis,⁶⁶ the most debilitating clinical sequela of this disorder.

Discussion

Over the past decade, an explosion of evidence has reinforced the neurogenic etiology of the CCDDs. The identification of specific gene mutations as the pathogenesis for these disorders has increased our knowledge of the importance of the normal protein products of these genes in the normal development of ocular cranial motor neurons and guidance of their axons to the appropriate end muscle targets. The combination of clinical findings, MR imaging, and gene product characterization has led to a greater understanding of the cascade of events that leads ultimately to the clinical presentation of these patients. This cascade starts as a gene mutation coding for an abnormal protein that leads to changes in the normal development of the nervous system that then results in structural abnormalities that culminate in anatomical dysfunction. The effort to gather this knowledge has spanned continents and has involved much collaboration among institutions.

Despite the advances in our understanding, much remains to be learned about these conditions. The variable expression pattern of patients with known mutations reveals the potential importance of additional genetic variation and environmental factors on the end phenotype. The high number of simplex CCDD patients with no identified gene mutation again highlights that environmental factors or somatic mutations may cause disruptions in development that result in a clinical presentation similar to that of underlying germline mutations. In many pedigrees there is little or no genetic data to point to the identity of the mutation. For these reasons, much work is still needed to obtain a better understanding of the environmental and genetic mechanisms that lead to dysinnervation in the CCDDs.

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FIGURE 1.

Motility of a patient with a KIF21A mutation. He also has aberrant innervation causing a Marcus Gun jaw wink. Reproduced with permission from Yamada, Hunter, et al., "A novel kif21a mutation in a patient with congenital fibrosis of the extraocular muscles and Marcus Gunn jaw-winking phenomenon," Arch Ophthalmol. 2005;123(9):1254–1259.

FIGURE 2.

(A–E) and (G, H) Eye manifestations of patients with CFEOM3 and TUBB3 mutations. (F) and (I) Concomitant deformities of the extremities in two patients. (J) MRI of the brainstem at the level of the oculomotor nerve. (K–L) MRI of posterior orbit of a patient with TUBB3 mutation (K) compared with a normal posterior orbit (L). Reproduced with permission from Tischfield MA, Baris HN, Wu C, et al., "Human TUBB3 mutations perturb microtubule dynamics, kinesin interactions, and axon guidance," Cell. Jan 8 2010;140(1):74–87. 153×207mm (300×300 DPI).

FIGURE 3.

Motility of a patient with horizontal gaze palsy and progressive scoliosis. MRI of spine revealing profound scoliosis below. Reproduced with permission from Jen JC, Chan WM, Bosley TM, et al., "Mutations in a human ROBO gene disrupt hindbrain axon pathway crossing and morphogenesis," Science. Jun 4 2004;304(5676):1509–1513.

FIGURE 4.

MRI studies comparing normal subjects (A–C) with patients with HGPPS (D–F) at similar anatomical levels. CC – corpus callosum, P - pons, M - medulla, * - fourth ventricle. Reproduced with permission from Jen JC, Chan WM, Bosley TM, et al., "Mutations in a human ROBO gene disrupt hindbrain axon pathway crossing and morphogenesis," Science. Jun 4 2004;304(5676):1509–1513. 368×235mm (72×72 DPI).

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Table 1

Summary of Disorders with Defined Genetic Basis Summary of Disorders with Defined Genetic Basis

Key: AD - autosomal dominant, AR - autosomal recessive, DS - Duane Syndrome, CFEOM - congenital fibrosis of extraocular muscles, HGPPS - horizontal gaze palsy with progressive scoliosis Key: AD – autosomal dominant, AR – autosomal recessive, DS – Duane Syndrome, CFEOM – congenital fibrosis of extraocular muscles, HGPPS – horizontal gaze palsy with progressive scoliosis

Table 2a

Quick-Look Differentiation of Syndromes with Similar Clinical Pictures for SALL4 Mutations Quick-Look Differentiation of Syndromes with Similar Clinical Pictures for *SALL4* Mutations

Table 2b

Quick-Look Differentiation of Syndromes with Similar Clinical Pictures for HOXA1 Mutations Quick-Look Differentiation of Syndromes with Similar Clinical Pictures for *HOXA1* Mutations

