

FIG. 5. Radiographs of Patient 3 compatible with the diagnosis of spondyloepipyseal dysplasia, Kozlowski type. A: Pelvis A-P at age 2 years. Persistent hypoplasia of the basilar portions of the iliac bones and round capital femoral epiphyses. The femoral necks are short, but have no overt metaphyseal abnormalities. B: Lateral spine at age 3 years. The vertebral bodies are flat with anterior wedging.

although there is large intra- and inter-familial variability [Dai et al., 2010a].

Patient 2 was diagnosed as having SEMD-M, but the mutation found, p799R, was identified previously in two cases of metatropic dysplasia by our group [Dai et al., 2010a]. The same TRPV4 mutation shows considerably different skeletal phenotypes in different patients [Dai et al., 2010b]. We reviewed the records of the cases with the p799R mutations, but found nothing to suggest the association of peripheral neuropathy. The neurological phenotype in Patient 2 was compatible with a motor and sensory axonal neuropathy predominant in the lower limb. Patient 3 had a novel TRPV4 mutation and was diagnosed as having SMD-K and a neurological phenotype compatible with scapulo-humeral spinal muscular atrophy. Thus, the three cases have different TRPV4 mutations and quite varied skeletal and neurological phenotypes.

Co-occurrence of skeletal and neurological phenotypes has been hinted at in a few instances. Chen et al. [2010] reported a threegeneration family with six patients harboring an \$542Y mutation. The patients in this family manifested as HMSN2C but in addition had moderate short stature. The diagnosis of the neuropathy was unequivocal, while characterization of skeletal changes was minimal and radiographic data were not presented. The description suggests to us that the family might have had brachyolmia or mild SMD-K. Zimon et al. [2010] reported an isolated Croatian patient (PN-1394.1 in the paper) who was diagnosed as having HMSN2, and also had scoliosis and short stature. Although the authors stated that skeletal abnormalities in the patient were too limited to make a formal diagnosis of a skeletal dysplasia, the radiographic description suggests brachyolmia. The patient had a de novo V620I mutation, which was previously shown to cause autosomal dominant brachvolmia [Rock et al., 2008; Dai et al., 2010a]. Recently, Unger et al. [2011] reported fetal akinesia as the presenting feature of severe metatropic dysplasia, which supported that certain *TRPV4* mutations can cause both skeletal and neuropathic phenotypes.

Earlier reports stressed that TRPV4 mutations associated with neuropathy involve substitutions of arginine residues at the ankyrin repeats [Auer-Grumbach et al., 2010; Chen et al., 2010; Deng et al., 2010; Landoure et al., 2010; Berciano et al., 2011; Klein et al., 2011]. However, two neuropathic mutations have been reported in the TRPV4-transmembrane domain (S542Y and V620I) [Chen et al., 2010; Zimon et al., 2010]. Neither of them is an arginine substitution. Conversely, many TRPV4 mutations identified in skeletal dysplasias are arginine substitutions and several are in the ankyrin repeats [Dai et al., 2010a]. Also, all mutations in our patients were not arginine substitutions, and the mutation in Patient 2 was not in the ankyrin repeat. Our conclusion from these findings is that genotype—phenotype association in TRPV4-pathies is not strict and may lead to clinical phenotypes manifesting with both skeletal and neurological abnormalities.

Pathogenesis remains enigmatic and contentious in TRPV4associated neuropathies, where some in vitro experiments indicated a possible loss-of-function for specific TRPV4 variants [Auer-Grumbach et al., 2010]. In contrast, all mutations reported to date, including the mutations associated with isolated neurological disease, seemingly activate non-selective Ca2+ channel function of TRPV4. Although recent in vitro evidence demonstrates that the extent of functional channel gain in Xenopus oocytes correlates apparently well with disease severity in the skeleton [Loukin et al., 2011], clinical evidence of eminent intra- and inter-familial variability strongly contradicts a simple pathogenic mechanism. Our findings prove that clinical variability associated with specific TRPV4 mutations may even extend to different organ systems. Understanding the molecular pathophysiology of pleiotropic TRPV4-pathies thus remains a challenge and awaits the generation of true disease models.

Thus, we have described clear-cut cases that show both skeletal dysplasia and axonal type peripheral neuropathy in association with TRPV4 mutations. It implies that the pathogenic mechanisms of phenotypes in the two systems are not mutually exclusive. As the developmental skeletal phenotype may be detected earlier than degenerative neurologic disorder, those patients confirmed to have skeletal dysplasias of TRPV4-pathy should be paid special attention to their neurologic abnormality. Any deficit developing in TRPV4-pathy skeletal dysplasia patients should be differentiated between TRPV4-related peripheral neuropathy versus cord or spinal nerve compression from vertebral abnormalities.

Despite the presence of unequivocal radiographic changes in all our patients, their heights were within the normal range. As short stature is often the primary indication of a skeletal dysplasia, it is possible that more patients with TRPV4-related peripheral neuropathy have skeletal manifestations, but have simply never been investigated because they are not short. Radiographic examination should be considered as a part of the standard work-up for patients compatible with TRPV4-related neuropathies. In clinical diagnosis and genetic counseling, the extensive phenotypic variability and reduced penetrance of neurologic phenotype of TRPV4-pathy should be taken into account.

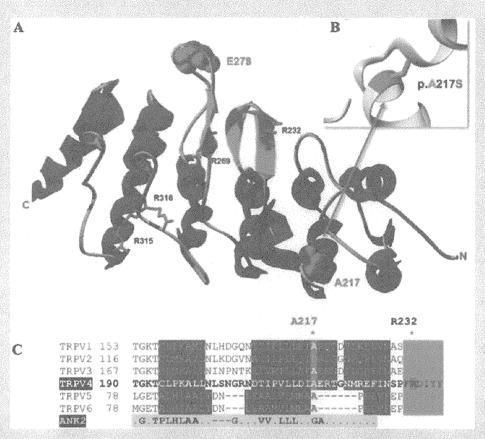


FIG. 6. A: A Ribbon diagram of the modeled ankyrin repeat region of human TRPV4. Residues E278 and A217 mutated in Patients 1 and 3, respectively, are depicted as red spheres, all other previously reported ankyrin repeat region residues affected by arginine substitutions in dominant neuropathies (R232, R269, R315, and R316) are represented as sticks. B: A close-up view of position 217 from a different angle. The non-polar to polar substitution p.A217S changes helical winding in the second ankyrin repeat region of TRPV4. The overlays of wild type alanine (green) and the substituted serine (red) are superposed in the ribbon and stick model to illustrate molecular dimensions of the respective amino acids. C: Alignment of the second ankyrin repeat region of human TRPVs, corresponding to amino acids 190–226 in TRPV4; ankyrin consensus residues conserved in TRPVs are included in yellow shading at the bottom. Blue shading indicates helical stretches confirmed in chicken Trpv4 and rat Trpv6 crystal structure, green shading indicates β-sheets; A217 (red) is conserved in all but one of TRPVs.

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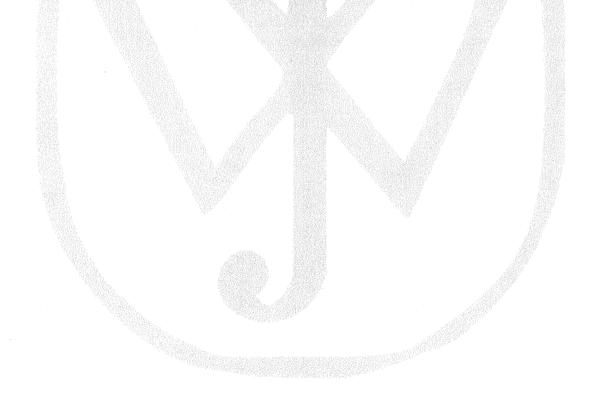
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Axial Spondylometaphyseal Dysplasia: Additional Reports

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Axial spondylometaphyseal dysplasia (SMD) (OMIM 602271) is an uncommon skeletal dysplasia characterized by metaphyseal changes of truncal-juxtatruncal bones, including the proximal femora, and retinal abnormalities. The disorder has not attracted much attention since initially reported; however, it has been included in the nosology of genetic skeletal disorders (Warman et al. (2011); Am J Med Genet Part A 155A:943-968] in part because of a recent publication of two additional cases [Isidor et al. (2010); Am J Med Genet Part A 152A:1550-1554]. We report here on the clinical and radiological manifestations in seven affected individuals from five families (three sporadic cases and two familial cases). Based on our observations and Isidor's report, the clinical and radiological hallmarks of axial SMD can be defined: The main clinical findings are postnatal growth failure, rhizomelic short stature in early childhood evolving into short trunk in late childhood, and thoracic hypoplasia that may cause mild to moderate respiratory problems in the neonatal period and later susceptibility to airway infection, Impaired visual acuity comes to medical attention in early life and function rapidly deteriorates. Retinal changes are diagnosed as retinitis pigmentosa or pigmentary retinal degeneration on fundoscopic examination and cone-rod dystrophy on electroretinogram. The radiological hallmarks include short ribs with flared, cupped anterior ends, mild spondylar dysplasia, lacy iliac crests, and metaphyseal irregularities essentially confined to the proximal femora. Equally affected sibling pairs of opposite gender and

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parental consanguinity are strongly suggestive of autosomal recessive inheritance, % 2011 Wiley-Liss, Inc.

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INTRODUCTION

The term spondylometaphyseal dysplasia (SMD) encompasses a heterogeneous group of disorders characterized by dysplastic changes in the metaphyses of tubular bones and metaphyseal equivalents of the spine and flat bones. Amongst the SMDs, the most common is SMD Kozlowski type (OMIM 184252), followed by SMD Sutcliffe (corner fracture) type (OMIM 184255), but there are also other rare types [Wirth, 2008]. Aside from SMD Kozlowski type, which is caused by heterozygous mutations in the transient receptor potential cation channel, subfamily V, member 4 gene (TRPV4) [Nishimura et al., 2010], the etiologies of the SMDs remain unknown. It is intriguing that certain types of SMDs present as a multi-system disorder, as exemplified by SMD with cone-rod dystrophy (OMIM 608940) [Sousa et al., 2008; Turell et al., 2010].

We previously reported on a new type of SMD, based on the clinical and radiologic observations in three children (a Japanese girl and two Korean siblings) [Ehara et al., 1997]. The disorder was termed "axial SMD", because the metaphyseal changes were confined to the truncal and juxtatruncal bones. The disorder was seen in association with ocular abnormalities, including retinitis pigmentosa (RP) and/or optic atrophy. Very recently, two additional cases have been reported confirming the axial SMD as a distinct entity [Isidor et al., 2010]. The Nosology Group of the International Skeletal Dysplasia Society included the disorder in "Nosology and classification of genetic skeletal disorders: 2010 version", and the group termed the entity SMD with retinal degeneration, axial type (OMIM 602271) [Warman et al., 2011]. With the addition of five newly identified cases (three sporadic patients and two siblings) and follow-up our previously reported sibling case, we are able to delineate the key clinical and radiographic features of this condition and hopefully facilitate the diagnosis of further individuals.

CLINICAL REPORTS

Patient 1

Patient 1 is a Japanese girl born to healthy, nonconsanguineous parents. Birth length was $47.6 \,\mathrm{cm}\,(-1.0\,\mathrm{SD})$. At birth, she had mild respiratory distress and a narrow thorax was noted. A tentative radiological diagnosis of Shwachman-Diamond syndrome was made. Ophthalmological screening at age 3 weeks disclosed RP (Fig. 1A), and no electrical activities were detectable by electroretinogram (ERG) at 4 months (Fig. 1B). Length at age 11 months was $64.3 \,\mathrm{cm}\,(-3.4\,\mathrm{SD})$.

Patient 2 and 3 (siblings)

Patients 2 and 3 have consanguineous parents (first cousins) of Saudi Arabian origin who also have four healthy children.

Patient 2, a girl, was born by vaginal delivery at 40 weeks' gestation after an unremarkable pregnancy. Apgar scores were 9 and 10 at 1 and 5 min, respectively. Birth length was 51 cm, and weight 3,440 g (both at 50th centile). At age 4 months, she was noted

to have micromelic short stature (<3rd centile) with rhizomelic shortening of the upper limbs and a narrow thorax with Harrison's grooves. At age 4 $^{7/12}$ years, height was 93.5 cm (-4.0 SD), and weight was 13.1 kg (-3.0 SD). Ophthalmological examination at age 1 year showed early signs of RP. She underwent ERG examinations twice at age 3 and 4 years, which showed no response to light flash stimulation indicative of advanced retinal dysfunction, and visually evoked response study showed delayed P100. At last examination, she had low visual acuity and no night vision but she was able to walk independently. Her visual acuity has deteriorated faster than her brother's.

Patient 3, the older brother of Patient 2, was born by vaginal delivery at 40 weeks' gestation after an unremarkable pregnancy. Apgar scores were 9 at 1 and at 5 min. Birth weight was 3,480 g, length was 51 cm, and OFC was 35 cm (all at 50th centile). He sat at age 6 months and walked at 10 months. His development was unremarkable except for visual dysfunction. He was noted to have RP at age 9 months, Mild hyperopia and astigmatism were found bilaterally at age 3 years. He had ERG three times at age 2 9/12 years, 6 years, and 7 years, all of which showed abnormal response to light flash stimulation. However, visually evoked response study was not significantly affected. At age 3 7/12 years, he was referred for genetic consultation because of failure to thrive, short stature, small chest, and visual problems. He showed micromelic short stature (<3rd centile) with rhizomelic shortening of the upper limbs and a narrow thorax with Harrison grooves. At age 7 years, height was 106.5 cm (-5.5 SD), weight was 18.5 kg (-2.5 SD), and OFC was 52.6 cm (-2.5 SD).

At last review, he had low visual acuity and no night vision but he was able to walk independently and could read with difficulty during daytime.

Patient 4

Patient 4 is a Korean boy, who was the second child of non-consanguineous, healthy parents. He came to medical attention at age 5 years because of short stature; $102.7 \, \text{cm} \, (-1.6 \, \text{SD})$. He had mild bowlegs and pectus excavatum. At that time, inward gaze of the eyes and impaired visual acuity were also noted. At age 6 years, mild RP was diagnosed on fundoscopic examination. Then, visual acuity progressively worsened. At age 9 years, vision was severely impaired (V.d. = 0.06; V.s. = 0.125). On last examination, at $10 \, \text{years}$ of age, he had proportionate short stature with a height of $125.8 \, \text{cm} \, (-2.0 \, \text{SD})$. In addition, modest narrow thorax, mild thoracic scoliosis, and rhizomelic shortening of the upper limbs were seen (Fig. 2A).

Patient 5

Patient 5 is a Japanese boy born to healthy, nonconsanguineous parents. He was delivered at 41 weeks' gestation. Birth length was $48.5 \,\mathrm{cm}$ ($-0.9 \,\mathrm{SD}$), weight was $3,052 \,\mathrm{g}$ ($-0.2 \,\mathrm{SD}$), and OFC was $34.0 \,\mathrm{cm}$ ($+0.1 \,\mathrm{SD}$). A narrow thorax was noted at birth. Apgar scores were 7 at 1 min and 5 at 5 min. He had moderate respiratory distress with laryngomalacia necessitating oxygen therapy in the neonatal period. The laryngomalacia improved gradually over the course of the first year. He was given a radiological diagnosis of

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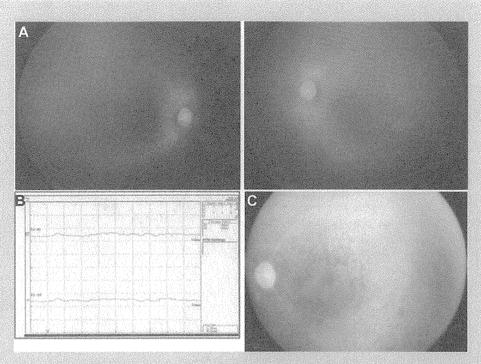


FIG. 1. Ophthalmological findings. A: in Patient 1, fundoscopy showed RP at age 2 months. B: in Patient 1, ERG did not trace electric activities at age 4 months. C: in Patient 5, fundoscopic findings were advanced RP with reduced retinal blood flow and optic nerve atrophy at age 12 years.

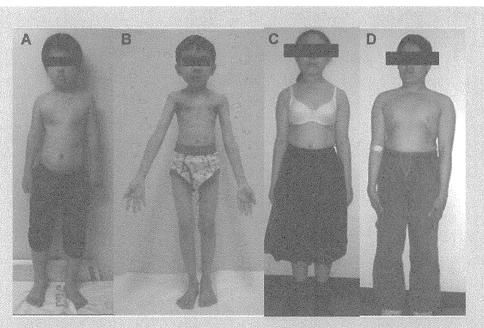


FIG. 2. Clinical photographs, A: Patient 4 at age 10 years. A mild narrow thorax and rhizomelic shortening of the upper limbs are noted. B: Patient 5 at age 12 years. Short trunk is striking with a mild narrow thorax. C: Patient 6 at age 15 years, (D) Patient 7 at age 23 years. Mild short trunk is evident, and the thorax is mildly narrow.

Jeune asphyxiating thoracic dysplasia in infancy. He had recurrent episodes of pneumonia until age 5 years. At age 6 years, severe short stature was recorded (-6.5 SD). Visual problems were suspected at an early age as he never showed any light response or the ability to track moving objects. Nystagmus was identified in early infancy. He was diagnosed as having macular RP at age 2 months. Visual evoked potential test showed no response. Severe hyperopia with disturbed visual acuity (V.d. = 0.01, V.s. = 0.01) was documented at age 5 years. Visual acuity gradually declined. He suffered from bilateral cataracts of the posterior subcapsular lense at 11 years. At age 12 years, fundoscopic findings showed advanced RP with reduced retinal blood flow and optic nerve atrophy (Fig. 1C). Clinical examination at that age demonstrated a height of 107.2 cm (-6.2 SD), arm span of 111.7 cm, and upper segment of 51 cm. He had a mild narrow thorax, mild scoliosis, rhizomelic shortening of the limbs, and markedly short trunk (Fig. 2D). Some permanent teeth had not yet erupted. Pulmonary function tests showed restrictive impairment with 38% forced vital capacity and 114.5% forced expiratory volume in 1 sec.

Patient 6 & 7 (siblings)

Patients 6 and 7 are Korean siblings, whose manifestations in childhood were previously reported [Ehara et al., 1997]. They were born to healthy, nonconsanguineous parents. Birth weight was normal. Short stature was noted in early childhood, as well as thoracic hypoplasia with susceptibility to airway infections. Height was 76 cm $(-5.5 \, \mathrm{SD})$ at age 3 years in the younger sister, and 108 cm $(-5.1 \, \mathrm{SD})$ at age 10 years in the older brother. The younger sister was diagnosed as having optic atrophy with nystagmus at age 3 years. Impaired visual acuity of the older brother came to attention

at age 6 months, and he was diagnosed as having optic atrophy and retinal degeneration associated with nystagmus at age 8 years. At the most recent examination, the sister was 15 years, and the brother was 23 years old. Their heights were 131 cm (-5.1 SD) and 144 cm (-4.9 SD), respectively. They presented with a narrow thorax, short-trunk, and rhizomelic shortening of the upper limbs (Fig. 2C,D). They were functionally blind at that time.

RADIOLOGICAL FINDINGS

The radiographic findings in all patients were similar but with a variable degree of severity. The phenotype evolved with age.

Chest

Short ribs with flared, cupped anterior ends were evident in the neonatal period (Fig. 3A). This finding became prominent in childhood, most striking in late childhood (Fig. 3B–I), and then less conspicuous in adolescence and adulthood (Fig. 3J,K).

Spine

Spondylar changes were mild in infancy and early childhood (Fig. 4A–D). Platyspondyly became more apparent in late childhood (Fig. 4E–H). Vertebral height increased in adolescence and normalized in adulthood (Fig. 4I,J).

Pelvis

Lacy ilia were discernible in the neonatal period and became overt in childhood. Metaphyseal irregularities of the proximal femora became manifest in infancy and then progressed. Coxa vara devel-

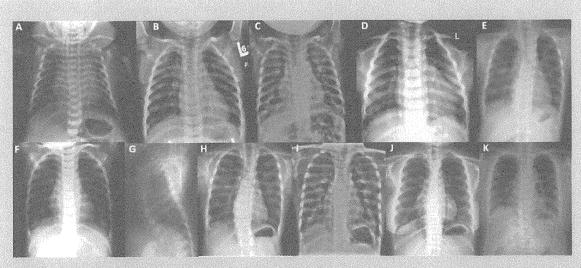


FIG. 3. Radiographs of the chest. A: Patient 1 at age 2 days, (B) Patient 2 at age 1^{4/12} years, (C) Patient 5 at age 2 years, (D) Patient 3 at age 4 years, (E) Patient 4 at age 5 years, (F) and (G) Patient 6 at age 8 years. Note a narrowthorax in (A), (C—E), and short ribs with cupped, flared anterior ends in all. Thoracic narrowing is modest in (B) and (F). Mild irregularities of the proximal humeral metaphyses are seen in (A,B) and (C). H: Patient 4 at age 10 years, (1) Patient 5 at age 12 years. Cupping and flaring of the anterior ends of the ribs are most conspicuous in late childhood. Thoracic narrowing is modest. J: Patient 6 at age 15 years, (K) Patient 7 at 23 age years. A narrow thorax is persistent, but cupping and flaring of the anterior ends of the ribs are less conspicuous than those at younger ages.

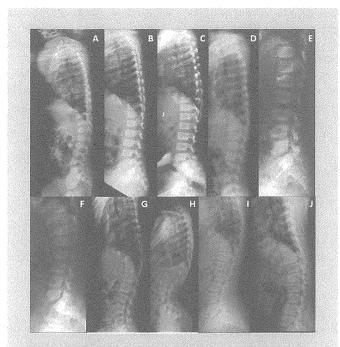


FIG. 4. Radiographs of spine. A: Patient 1 at age 6 months. The vertebral bodies are somewhat ovoid, but platyspondyly is not evident. B: Patient 2 at age 1 4/12 years, (C) Patient 3 at age 4 years, (D) Patient 4 at age 5 years. Mild platyspondyly is evident. E: Patient 6 at 8 age years, (F) Patient 7 at age 10 years, (G) Patient 4 at age 10 years, (H) Patient 5 at age 12 years. Platyspondyly is most conspicuous in late childhood. I: Patient 6 at age 15 years, (J) Patient 7 at age 23 years. Platyspondyly is discernible but milder than that in childhood.

oped in late childhood (Fig. 5A–I). The iliac and metaphyseal changes diminished during adolescence, leaving only coxa vara (Fig. 5J,K).

Limbs

The proximal humeri showed mild metaphyseal irregularities (Fig. 3 A–C). Metaphyseal changes were absent or very mild in the knee (Fig. 6A–G).

Hands

Hands were unremarkable in all patients (data not shown).

DISCUSSION

Based on our experiences and the observations reported by Isidor et al. [2010], it is clear that axial SMD is a distinctive disease entity with recognizable clinical and radiographic features. "A new form of oculoskeletal syndrome" reported by Megarbane et al. [2004] may represent the same disorder. The clinical manifestations of the present and previously reported patients are summarized in Table I. The clinical hallmarks include postnatal growth deficiency, thoracic

hypoplasia, and retinal abnormalities. Equally affected siblings of opposite gender and the presence of consanguinity in some parents are strongly suggestive of an autosomal recessive pattern of inheritance.

Although birth length is in the normal range, short stature with rhizomelic limb shortening becomes apparent during childhood. Short stature is mild to moderate during childhood; however, growth failure is progressive, and final height may be less than —5 SD. Progressive shortening of the trunk over time results in the ultimately short-trunk body proportion. Thoracic hypoplasia with mild to moderate respiratory distress in the neonatal period is apparent in some cases, while it may be asymptomatic in others. The narrow thorax occasionally gives rise to Harrison grooves and susceptibility to airway infection in infancy and early childhood. Laryngotracheomalacia may contribute to the respiratory problems.

RP or pigmentary retinal degeneration is detectable during childhood and retinal changes may even be observed in the neonatal period. Electroretinography reveals cone-rod dystrophy [Isidor et al., 2010]. Secondary optic atrophy may ensue, and one child (Patient 5) presented cataracts that might be secondary to RP [Jackson et al., 2001]. The prognosis for vision is unfavorable.

As discussed by Isidor et al. [2010], the differential diagnosis Shwachman-Bodian-Diamond syndrome (OMIM 260400), Jeune asphyxiating thoracic dysplasia (OMIM 208500), Saldino-Mainzer syndrome (OMIM 266920), Dyggve-Melchior-Clausen (DMC) dysplasia (OMIM 223800), and SMD with conerod dystrophy (OMIM 608940). Shwachman-Diamond syndrome causes thoracic hypoplasia and metaphyseal dysplasia most conspicuously in the proximal femora. However, Shwachman-Bodian-Diamond syndrome, unlike axial SMD, is associated with neutropenia and pancreatic exocrine dysfunction but not retinal and spondylar changes. Both Jeune asphyxiating thoracic dysplasia and Saldino-Mainzer syndrome manifest thoracic hypoplasia and retinopathy. Nevertheless, progressive nephropathy and brachydactyly are seen in these disorders and are conspicuously absent in axial SMD. Lacy ilia and spondylar dysplasia in axial SMD may raise a suspicion of DMC. However, DMC shows more severe platyspondyly and epimetaphyseal dysplasia but not retinal changes. SMD with cone-rod dystrophy is a recently identified skeletal dysplasia associated with retinal cone-rod dystrophy. The clinical and radiological pattern in SMD with cone rod dystrophy is similar to that of axial SMD. However, there are clinical and radiological differences between both disorders. Visual impairment is milder in SMD with cone rod dystrophy. Affected individuals do not show complete loss of visual acuity. On the other hand, the skeletal changes of SMD with cone rod dystrophy are much more severe than those in axial SMD. Generalized metaphyseal dysplasia and more severe platyspondyly in SMD with cone rod dystrophy contrast with metaphyseal dysplasia confined to the juxtatruncal bones and mild platyspondyly in axial SMD.

In the neonatal period, axial SMD should be differentiated from SMD Sedaghatian type (OMIM 250220), a perinatally lethal osteochondrodysplasia comprising minor facial, cardiac and cerebral anomalies [Elçioglu and Hall, 1998]. Unlike axial SMD, SMD Sedaghatian type manifests overt metaphyseal dysplasia and lacy ilia in the neonatal period.

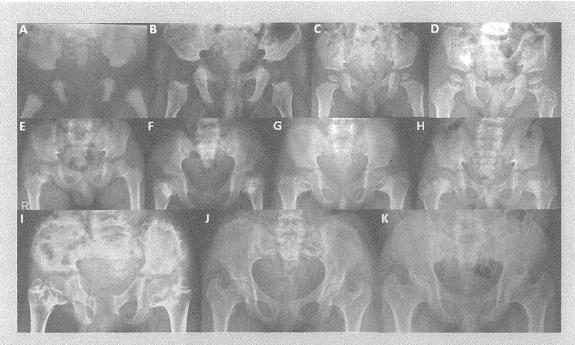


FIG. 5. Radiographs of pelvis. A: Patient 1 at 2 days. The ilia are somewhat hypoplastic but lacy ilia are not evident. B: Patient 1 at age 6 months. The iliac crests are somewhat irregular. C: Patient 2 at age 1 4/12 years. The iliac crests and proximal femoral metaphyses are irregular. D: Patient 3 at age 4 years, (E) Patient 4 at age 5 years. Lacy ilia and metaphyseal irregularities of the proximal femora are apparent. F: Patient 6 at age 8 years, (G) Patient 7 at age 10 years, (H) Patient 4 at age 10 years, (I) Patient 5 at age 12 years. Proximal femoral metaphyseal irregularities with coxa vara are conspicuous in all patients. Lacy ilia are varied among the patients. J: Patient 6 at age 15 years, (K) Patient 7 at age 23 years. Lacy ilia and metaphyseal irregularities already diminish, but coxa vara is persistent.

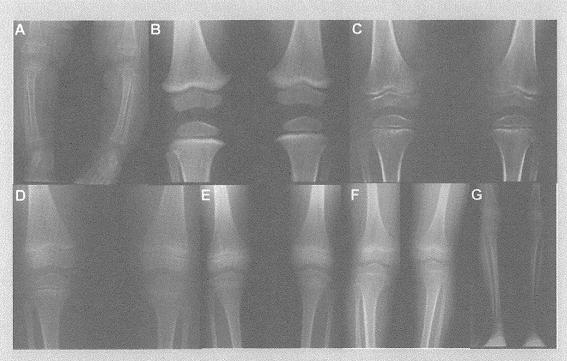


FIG. 6. Radiographs of lower limbs. A: Patient 1 at age 6 months, (B) Patient 2 at age 1 4/12 years, (C) Patient 3 at age 4 years, (D) Patient 6 at age 8 years, (E) Patient 7 at age 10 years, (F) Patient 4 at 10 years, (G) Patient 5 at 12 years. Metaphyseal changes are absent or very mild in the knee but the knee epiphyses are slightly flattened, notably in (B) and (C).

TABLE I. Clinical Characteristics of Patients With Axial SMD Isidor et al. Megarbane et al. Ehara et al. [2010] Present patients [1997] [2004] 2 3 5 6* 78 1 4 Patient Case1 Patient 1 Patient 2 Single case Race Japanese Saudi Arabian Saudi Arabian NA Lebanese Korean Japanese Korean Korean Japanese NA Sex Female Female Male Male Male Female Male Female Male Male Female Age at last visit (urs) 0.9 4.5 7 10 12 23 5 15 14 15 4 Family history Sporadic Sib Sib Sporadic Sporadic Sib Sib Sporadic Sporadic Sporadic Sporadic Short stature -3.4 SD -4.0 SD -5.5 SD -2.0 SD -6.2 SD-5.1 SD -4.7 SD -3.6 SD -3.2 SD -3.6 SD<3% tile Small thorax + Short trunk NA NA NA Rhizomelic shortness + + Respiratory disturbance at birth NA. NA NA NA NA NA Respiratory infection in childhood NA NA NA NA NA NA NA RP/optic atrophy + + 6^b Age of Dx of RP (years) 0 0.9 0.1 8^c 0.1 6.5

+

NA, not available; RP, retinitis pigmentosa.

Scoliosis

^{*}Follow-up study of cases previously described as Case 3 and 2, respectively by Ehara et al. [1997].

bimpaired visual acuity was noticed at 5 years of age.

Impaired visual aculty was noticed at 6 months of age.

The molecular basis of axial SMD remains elusive; however, homozygosity mapping and whole genome sequencing techniques should elucidate the disease-causing gene in the near future. Further case reports, sample registration, and investigations will be invaluable in order to thoroughly understand this entity.

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ORIGINAL ARTICLE

A founder mutation of CANT1 common in Korean and Japanese Desbuquois dysplasia

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Desbuquois dysplasia (DBQD) is a severe skeletal dysplasia of autosomal recessive inheritance. DBQD is classified into types 1 and 2 based on presence or absence of hand anomalies. In a previous study, we found a CANT1 (for calcium-activated nucleotidase 1) mutation, c.676G > A in five DBQD families. They were all East Asians (Japanese or Korean). The high prevalence of the same mutation among Japanese and Korean suggested that it is a common founder mutation in the two populations. To examine a possible common founder, we examined the region around CANT1 in chromosomes with c.676G>A mutation by genotyping polymorphic markers in the region for the families. We examined their haplotypes using the family data. We identified in all families a common haplotype containing the CANT1 mutation that ranged up to 550 kb. The two unrelated carriers of the mutation in general populations in Korea and Japan could also have the haplotype. We estimated the age of the founder mutation as \sim 1420 years (95% CI=880–1940 years). The c.676G > A mutation of CANT1 commonly seen in Japanese and Korean DBQD should be derived from a common founder.

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Keywords: calcium-activated nucleotidase 1; common founder; common haplotype; desbuquois dysplasia

INTRODUCTION

Desbuquois dysplasia (DBQD; MIM #251450) is a severe skeletal dysplasia of autosomal recessive inheritance that belongs to 'multiple dislocation group? 1 DBQD is clinically heterogeneous and classified into two types on the basis of presence (type 1) or absence (type 2) of characteristic hand anomalies, which consist of an extra ossification center distal to the second metacarpal, delta phalanx, bifid distal thumb phalanx and dislocation of the inter-phalangeal joints.² Type 2 DBQD contained a further clinical subtype, Kim variant, which is characterized by short metacarpals with elongated phalanges that result in nearly normal length of the fingers.

Recently, Huber et al.4 identified mutations in the gene encoding the calcium-activated nucleotidase 1 (CANT1) in DBOD type 1, which is followed by Faden et al.5 We also found CANT1 mutations in DBQD type 2 and Kim variant. In our series, we identified CANT1 mutations in all seven patients from five unrelated families with DBQD Kim variant, and all patients had c.676G>A (p.V226M).

The five families were Japanese or Korean. c.676G>A was also found in 1/754 Japanese and 1/187 Korean controls in a heterozygous state.6 These results led us consider that the mutation may be inherited from a common founder(s) among Japanese and Korean. To test the hypotheisis, we examined haplotypes of the patients around CANTI and found that the mutation was on a common haplotype background of $\sim 500 \, \text{kb}$.

MATERIALS AND METHODS

Subjects

Seven DBQD patients from five families (four Japanese and three Korean) and their parents were examined. One Japanese family (DB1) was consanguineous (1st cousin marriage) and two sib patients from the family had c.676G > A in a homozygous state. The other patients were of non-consanguineous and compound heterozygotes of the mutations.⁶ Their phenotypes were all diagnosed as DBQD Kim variant as previously reported.³ One Japanese (CT1) and one Korean (CT2) subject from general populations who were found to have c.676G > A in a heterozygous state in the previous study were also examined.3

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Genotyping

Genomic DNA was extracted from blood by standard procedures or from saliva using Oragene DNA Self-Collection kit (DNA Genotek, Ottawa, Ontario, Canada). Affymetrix 10K Gene-Chip microarray (Affymetrix, Santa Clara, CA, USA) was used for family DB1 to determine homozygous regions containing CANT1 inherited from a common ancestor in the consanguineous family. Single nucleotide polymorphisms (SNPs) for the founder haplotype determination were genotyped by direct sequencing. The regions containing targeted SNPs were amplified by PCR from genomic DNA, and then PCR products were diluted and sequenced by using an ABI Prism 3730 automated sequencer (PE Biosystems, Tokyo, Japan). We separated 5'-FAM—labeled PCR products containing microsatellites by sizes on an ABI Prism 3730 automated sequencer (PE Biosystems) against the Genescan-500LIZ size standard. PCR primers for the polymorphisms are listed in Supplementary Table 1.

Haplotype analysis

We determined the mutation-containing haplotype of patients according to the genotypes of the trio (the patient and parents). We determined the homozygous region of DB1 using the microarray data of the family members. All the patients were genotyped for the markers selected from the homozygous region of DB1. The genotyping of the makers for their parents were examined only when the patients were heterozygotes. The boundaries of the common haplotype were determined by sequential genotyping of SNPs in flanking regions. The SNPs were selected from the HapMap database (Phase II+ III, release 27).

Estimation of the mutation age

We estimated the age of the founder mutation by the method previously described.^{7,8} We used the following equation:

$$g = \log[(p_d - p_n)/(1 - p_n)]/\log(1 - \theta),$$

where p_d and p_n denote frequencies of the ancestral allele at the marker locus on the chromosomes with A and G alleles at the disease locus. θ denotes the recombination fraction between the marker and the disease locus. The age of the mutation was corrected by taking the population growth into account using the following equations:

$$g_{c}=g+g_{0},$$

$$g_0 = -(1/d)\ln[\theta \times e^d/(e^d - 1)],$$

where d denotes the population growth rate. Physical and genetic map positions in bp and cM for SNP loci were obtained from the HapMap database (Phase II, release 21).

RESULTS

Genotyping and identification of a common haplotype

Using the SNP microarray data, we first determined a homozygous region of the genome that contained CANT1 in the consanguineous family, DB1. The family had a stretch of SNPs that was homozygous in the patients and heterozygous in the parents. The homozygous stretch flanked by rs2934226 and rs2306755 extended up to ~9.8 Mb. Of 48 SNPs that composed the stretch in the microarray data, we selected 10 SNPs around CANT1; six were at the 3' to CANT1 and four were at the 5' to CANT1. We genotyped them for the five families. A possible common haplotype was restricted in the region between rs6501224 and rs2377402.

To confirm the common haplotype and define its more detailed boundary, we selected 13 SNPs in the boundary regions from a database and genotyped them for the five families. In the SNPs we could get a definite haplotype result, all in a region from rs10512617 to rs2045660 showed a common haplotype. The nearest SNPs to rs10512617 and rs2045660 that defined the different haplotypes in any of the five families were rs11077391 and rs2377309, respectively. At the SNPs between

Table 1 Haplotypes of markers with the c.676G>A mutation in the Desbuquois dysplasia (DBQD) families and control subjects

		Subject							
			DB	QD Fat	Control				
Marker	Distance ^a	DB1	DB2	DB3	DB4	DB5	CT1	СТ2	
rs7405591	1712736	С	Т	Т	Т	С	τ	Т	
rs4789523	978 405	T	T	T/Cb	T	T	T/C	T/C	
rs4129767	587 275	С	С	С	С	T	С	C/T	
rs6501224	488 804	G	G	С	G	G	G	G	
rs16971539	434 643	Α	G	G/Ab	G	G	G	G	
rs9899295	402 575	Α	G	Α	G	G	A/G	G	
rs9896451	347 233	G	G	Α	G	G	G	G/A	
rs11077391	330 052	G	Α	Α	G	Α	G	G	
rs10512617	297 708	G	G	G	G	G	G	G	
rs4103047	254 069	С	С	С	C	С	С	C/G	
rs3744801	197 531	С	С	C	С	С	C/T	С	
rs2889529	93 784	C	С	С	С	С	C/T	C/T	
c.676G>A	0								
rs8077024	2 287	С	С	С	С	С	С/Т	С	
D17S1847	33 602	185	185	185	185	185	185/191	185/189	
rs2707047	132 355	G	G	G	G	G	G/A	G/A	
rs2612787	132449	T	T	T	T	T	T/C	T/C	
rs9302889	168892	T	T	T	T	T	T/A	T	
rs12600665	180 703	G	G	G	G	G	G/A	G	
rs2045660	185 092	С	С	С	С	С	С	C	
rs2377309	201 821	С	С	С	С	C/Tb	С	C/T	
rs2612753	220795	G	G	G	G	T	G/T	G	
rs1000791	274619	T	Α	T	T	T	T	T/C	
rs2377402	340 187	Α	С	С	Α	С	С	С	

^aBase number between the marker and the disease locus. ^bEquivocal because of the double heterozygosity of the parents.

rs10512617 and rs2045660, the two carriers of the mutation (CT1 and CT2) had all the alleles that composed the common haplotype. Therefore, they could also have the same haplotype (Table 1).

Estimation of the mutation age

The markers that defined the haplotype of the families were used for estimation of the mutation age. The disease chromosomes of DB1 were counted as one because of its consanguinity. pd was calculated in five chromosomes with A allele at the disease locus from the five DBQD families. As frequency of the mutation (A allele) was quite low in control subjects, we regarded the allele frequency of markers in the general Japanese population as p_n , and the allele frequencies were obtained from NCBI dbSNP database. The population growth rate, d was set to 0.08 as previously described.7 The allele, frequency of which was higher in the five chromosomes than that in the general population was regarded as the ancestral allele. We excluded the markers in the region less than 60 kb from the disease locus. The estimated ages by different markers ranged from 30 to 246 generations (mean=71; 95% CI=44-97; median=41; standard deviation=58) (Table 2). If we assumed an intergenerational time of 20 years, the age of the mutation was estimated to be 1420 years (95% CI=880-1940 years; median=820 years; standard deviation=1160 years).

DISCUSSION

We have demonstrated that the c.676G>A mutation in Japanese and Korean had a common founder because all five chromosomes with

Table 2 Estimated age of the c.676G > A mutation

Marker	Pd	Pn	θ	g	gO	gc
rs7405591	0.4	0.295	0.0545	34.0	4.3	38.3
rs4129767	0.8	0.488	0.0191	25.7	17.4	43.1
rs6501224	8.0	0.698	0.0145	74.5	20.9	95.4
rs9899295	0.6	0.430	0.00998	120.6	25.5	146.1
rs9896451	8.0	0.488	0.00432	114.3	36.0	150.3
rs11077391	0.6	0.295	0.00401	208.6	37.0	245.6
rs10512617	1	0.400	0.00394	0	37.1	37.1
rs4103047	1	0.659	0.00391	0	37.2	37.2
rs3744801	1	0.523	0.00385	0	37.4	37.4
rs2889529	1	0.233	0.00187	0	46.5	46.5
c.676G>A			0			
rs2707047	1	0.145	0.00407	0	36.7	36.7
rs2612787	1	0.024	0.00408	0	36.7	36.7
rs9302889	1	0.727	0.00691	0	30.1	30.1
rs12600665	1	0.500	0.00722	0	29.6	29.6
rs2045660	1	0.576	0.00727	0	29.5	29.5
rs2612753	8.0	0.589	0.0106	62.8	24.8	87.6
rs1000791	0.8	0.625	0.0159	47.6	19.7	67.3
rs2377402	0.4	0.157	0.0201	61.3	16.8	78.1

c.676A in DBQD patients had a common haplotype composed of nine markers around this disease locus. In addition, the genotype of the two unrelated normal subjects with this mutation contained all the alleles that composed the common haplotype, suggesting that the founder mutation has widely spread among Korean and Japanese populations. The haplotype region ranged 480-550 kb. We estimated the age of this mutation at ~1400 years (95% CI=880-1940 years). Although the calculated CI is quite wide, we can estimate the c.676G>A mutation dates back to a time around the late Kofun era. Because the allele frequency of c.676G>A in Korean controls is much higher than that in Japanese controls, we speculated that the common founder was a Korean, and this mutation spread from Korea to Japan. This speculation is not contradictory to migration from Korea to Japan in the ancient history.

Although the exact prevalence of DBQD is difficult to determine, DBQD is generally considered as a very rare disease. However, the carrier frequency of the common mutation is considerably high in Korean. From our experience,6 the frequency of the CANT1 mutation is speculated as about double of that of the common mutation. Thereafter, prevalence of DBQD is $\sim 1/35000$ in Korean and \sim 1/600 000 in Japanese; the former is far higher than our impression in daily practice. Many DBQD patients caused by CANT1 mutations may be pre- or peri-natally lethal, although all the East Asian cases of DBQD with CANT1 mutations so far reported had good prognosis. Alternatively, many DBQD may be undiagnosed and/or be put into the waste box of diagnosis as unknown 'multiple dislocation group' of skeletal dysplasia. Diagnosis of DBQD based on clinical and/or radiographic information is sometimes very difficult because of the overlapping phenotype with other diseases.10

In DBQD patients, the founder mutation was identified in all the seven Kim variant patients, and only in the Kim variant patients to our knowledge. Therefore, this mutation may be closely related with specific phenotypes of the Kim variant; that is, accelerated carpal bone ages in childhood, short metacarpals, elongated appearance of phalanges and absence of accessory ossification center distal to the second metacarpal and thumb anomalies.3 Further accumulation of Korean and Japanese DBQD patients and examination of their CANT1 mutations are necessary to conclude relation of the founder mutation and the specific phenotype. The founder mutation may present with other phenotype. Moreover, further collection of DBQD patients and screening of CANT1 mutations in other populations would help confirming the phenotype-genotype relation. Other mutation (s) may present with the Kim-type DBQD phenotype. The diagnosis of DBQD is sometimes very difficult at present; however, we think identification of the specific phenotypes of hands followed by examination of the founder mutation in CANT1 would efficiently led us to definite diagnosis in Japanese and Korean patients suspected for having DBQD.

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Supplementary Information accompanies the paper on Journal of Human Genetics website (http://www.nature.com/jhg)

	Name of Disorder	疾患名	遗伝子*	ONJ	Gendia	GeneTests	国内コマーシャルベース	研究ベース
	FGFR3 group Thanatophoric dysplasia type 1 (TD1) Thanatophoric dysplasia type 2 (TD2) Severe achondroplasia with developmental delay and acanthosis nigricans (SADDAN) Achondroplasia Hypochondroplasia Hypochondroplasia	1. FGFR3グループ 致死性骨異形成症1型(TD1) 致死性骨異形成症2型(TD2) SADDAN(重症軟骨無形成症, 発達遅滞, 黒色表皮腫) 軟骨無形成症 軟骨低形成症 軟骨低形成症 軟骨低形成症検異形成症	FGFR3 FGFR3 FGFR3 FGFR3		00 0 00	0 0 0 0 0		可能(応相談) 可能(応相談) 可能(応相談) 可能(応相談) 可能(応相談) 可能(応相談)
	2. Type 2 collagen Group and similar disorders Achondrogenesis type 2 (ACG2; Langer-Saldino) Platyspondylic dysplasia, Torrance type Hypochondrogenesis Spondyloepiphyseal dysplasia congenital (SEDC) Spondyloepimetaphyseal dysplasia (SEMD) Strudwick type Kniest dysplasia Spondyloperipheral dysplasia Mild SED with premature onset arthrosis SED with metatarsal shortening (formerly Czech dysplasia) Stickler syndrome type 1	2. 2型コラーゲングループと類似疾患軟骨無発生症2型(ACG2: Langer-Saldino型)扁平椎異形成症, Torrance型軟骨低発生症先天性脊椎骨端異形成症(SEDC)脊椎骨端骨幹端異形成症(SEMD), Strudwick型Kniest骨異形成症脊椎末梢異形成症存性関節症を伴う軽症型脊椎骨端異形成症中足骨短縮を伴う脊椎骨端異形成症の足管短縮を伴う脊椎骨端異形成症(前Czech異形成Stickler症候群1型	COL2A1 COL2A1 COL2A1 COL2A1 COL2A1 COL2A1 COL2A1		0 0 0 0	0000000		可能(応応相報談) 可可能能(応応応応応応応応応応応応応応応応応応応応応応応応応応応応応応応応応応
— 103	Stickler-like syndrome(s) 3. Type 11 collagen Group Stickler syndrome type 2 Marshall syndrome Fibrochondrogenesis Otospondylomegaepiphyseal dysplasia (OSMED), recessive type Otospondylomegaepiphyseal dysplasia (OSMED), dominant type (Weissenbacher-Zweymuller syndrome, Stickler syndrome type 3)	Stickler様症候群 3. 11型コラーゲングループ Stickler症候群2型 Marshall症候群 線維性軟骨発生症 耳脊椎巨大骨端異形成症(OSMED), 劣性遺伝型 耳脊椎巨大骨端異形成症(OSMED), 優性遺伝型 (Weissenbacher-Zweymuller症候群3型)	COL11A1 COL11A1 COL11A1 COL11A2 COL11A2		00 0	00000		可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)
Existing and the second	4. Sulphation disorders group Achondrogenesis type 1B (ACG1B) Atelosteogenesis type 2 (AO2) Diastrophic dysplasia (DTD) MED, autosomal recessive type (rMED; EDM4) SEMD, PAPSS2 type Chondrodysplasia with congenital joint dislocations, CHST3 type (recessive Larsen syndrome) Ehlers-Danlos syndrome, CHST14 type ("musculo-skeletal variant")	4. 硫酸化障害グループ 軟骨無発生症1B型(ACG1B) 骨発生不全症2型(AO2) 捻曲性骨異形成症(DTD) MED, 常染色体劣性遺伝型(rMED, EDM4) SEMD PAPSS2型 先天性関節脱臼を伴う軟骨異形成症, CHST3型(劣性 Larsen症候群) Ehlers-Danlos症候群, CHST14型("筋骨格変異")	SLC26A2 (DTDST) SLC26A2 (DTDST) SLC26A2 (DTDST) SLC26A2 (DTDST) SLC26A2 (DTDST) PAPPSS2 CHST3 CHST14		0000	0000		可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)
	5. Perlecan group Dyssegmental dysplasia, Silverman-Handmaker type Dyssegmental dysplasia, Rolland-Desbuquois type Schwartz-Jampel syndrome (myotonic chondrodystrophy)	5. Perlecanグループ 分節異常骨異形成症, Silverman-Handmaker型 分節異常骨異形成症, Rolland-Desbuquois Schwartz-Jampel症候群(筋ミオトニー軟骨異栄養症)	HSPG2 HSPG2 HSPG2		000	0 0 0		可能(応相談) 可能(応相談) 可能(応相談)
	6. Aggrecan group SED (Spondyloepiphyseal Dysplasia), Kimberley type SEMD (Spondyloepimetaphyseal Dysplasia), Aggrecan type Familial osteochondritis dissecans	6. Aggrecanグループ SED, Kimberley型 SEMD, Aggrecan型 家族性離断性骨軟骨症	ACAN ACAN ACAN					可能(応相談) 可能(応相談) 可能(応相談)
	7. Filamin group and related disorders Frontometaphyseal dysplasia Osteodysplasty Melnick-Needles Otopalatodigital syndrome type 1 (OPD1) Otopalatodigital syndrome type 2 (OPD2) Terminal Osseous Dysplasia with Pigmentary Defects (TODPD) Atelosteogenesis type 1 (AO1) Atelosteogenesis type 3 (AO3)	7. Filaminグループと関連異常 全頭骨幹端異形成症 異形成骨症Melnick-Needles型 耳口蓋指症候群1型(OPD1) 耳口蓋指症候群2型(OPD2) 色素異常を伴う末端骨形成異常症(TODPD) 骨発生不全症1型(AO1) 骨発生不全症3型(AO3)	FLNA FLNA FLNA FLNA FLNA FLNB FLNB		0000 00	0000000		可能(応相談) 可能(応相談) 可能(応相談) 可能(応相談) 可能(応応相談) 可能(応応相談) 可能(応相談)

Larsen syndrome (dominant) Spondylo-carpal-tarsal dysplasia Spondylo-carpal-tarsal dysplasia Franck – ter Haar syndrome Serpentine fibula – polycystic kidney syndrome	Larsen症候群 (優性) 脊椎 • 手根骨 • 足根骨異形成症 脊椎 • 手根骨 • 足根骨異形成症 Franck – ter Haar症候群 蛇行腓骨 • 多囊胞腎症候群	FLNB FLNB FLNBと非連鎖	(可能(応相談) 可能(応相談)
8. TRPV4 group Metatropic dysplasia Spondyloepimetaphyseal dysplasia, Maroteaux type (Pseudo- Morquio syndrome type 2) Spondylometaphyseal dysplasia Kozlowski type Brachyolmia, autosomal dominant type Familial digital arthropathy with brachydactyly	TRPV4 group 変容性骨異形成症 脊椎骨端骨幹端異形成症, Maroteaux型 (偽性 Morquio症候群2型) 脊椎骨端異形成症Kozlowski型 短体幹症, 常染色体優性遺伝型 短指症を伴う家族性指関節症	TRPV4 TRPV4 TRPV4 TRPV4	(可能(応相談) 可能(応相談) 可能(応相談) 可能(応相談)
9. Short-rib dysplasias (with or without polydactyly) Group Chondroectodermal dysplasia (Ellis-van Creveld) SRP type 1/3 (Saldino-Noonan/Verma-Naumoff) SRP type 1/3 (Saldino-Noonan/Verma-Naumoff) SRP type 1/3 (Saldino-Noonan/Verma-Naumoff) SRP type 2 (Majewski) SRP type 4 (Beemer)	9. 短肋骨異形成症 (多指症を伴う/伴わない)グループ 軟骨外胚葉性異形成症 (Ellis-van Creveld) SRP 1/3型 (Saldino-Noonan/Verma-Naumoff) SRP 1/3型 (Saldino-Noonan/Verma-Naumoff) SRP 1/3型 (Saldino-Noonan/Verma-Naumoff) SRP 2型 (Majewski) SRP 4型 (Beemer)	, EVC, EVC2 DYNC2H1 IFT80 DYNC2H, IFT80と非連鎖 DYNC2H1	()	
Oral-Facial-Digital syndrome type 4 (Mohr-Majewski) Asphyxiating thoracic dysplasia (ATD; Jeune) Asphyxiating thoracic dysplasia (ATD; Jeune) Asphyxiating thoracic dysplasia (ATD; Jeune) Thoracolaryngopelvic dysplasia (Barnes)	口·顏面·指症候群4型 (Mohr-Majewski) 窒息性胸郭異形成症 (ATD; Jeune) 窒息性胸郭異形成症 (ATD; Jeune) 窒息性胸郭異形成症 (ATD; Jeune) 胸郭咽頭骨盤異形成症 (Barnes)	IFT80 DYNC2H1 DYNC2H, IFT80と非連鎖	C		
10. Multiple epiphyseal dysplasia and pseudoachondroplasia Group Pseudoachondroplasia (PSACH) Multiple epiphyseal dysplasia (MED) type 1 (EDM1) Multiple epiphyseal dysplasia (MED) type 2 (EDM2) Multiple epiphyseal dysplasia (MED) type 3 (EDM3) Multiple epiphyseal dysplasia (MED) type 5 (EDM5) Multiple epiphyseal dysplasia (MED) type 6 (EDM6) Multiple epiphyseal dysplasia (MED), other types Stickler syndrome, recessive type Familial hip dysplasia (Beukes) Multiple epiphyseal dysplasia with microcephaly and nystagmus (Lowry-Wood)	10. 多発性骨端異形成症および偽性軟骨無形成症グループ 偽性軟骨無形成症(PSACH) 多発性骨端異形成症(MED) 1型 (EDM1) 多発性骨端異形成症 (MED) 2型 (EDM2) 多発性骨端異形成症 (MED) 5型 (EDM3) 多発性骨端異形成症 (MED) 5型 (EDM6) 多発性骨端異形成症 (MED) 6型 (EDM6) 多発性骨端異形成症 (MED),他の型 Stickler症候群、劣性遺伝型 家族性臼蓋形成不全症 (Beukes) 小頭症と眼振を伴う多発性骨端異形成症 (Lowry- Wood)	COMP COMP COL9A2 COL9A3 MATN3 COL9A1	0		可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)可能(応相談)
11. Metaphyseal dysplasias Metaphyseal dysplasia, Schmid type (MCS) Cartilage-Hair-Hypoplasia (CHH; metaphyseal dysplasia, McKusick type) Metaphyseal dysplasia, Jansen type Eiken dysplasia Metaphyseal dysplasia with pancreatic insufficiency and cyclic neutropenia (Shwachman-Bodian-Diamond syndrome, SBDS) Metaphyseal anadysplasia type 1 Metaphyseal anadysplasia type 2 Metaphyseal dysplasia, Spahr type Metaphyseal Acroscyphodysplasia (various types) Genochondromatosis (type 1/type 2) Metaphyseal chondromatosis with D-2-hydroxyglutaric aciduria	11. 骨幹端異形成症 骨幹端異形成症、Schmid型(MCS) 軟骨・毛髪低形成症(CHH; 骨幹端異形成症, McKusick型) 骨幹端異形成症 Eiken異形成症 膵不全、周期性好中球減少を伴う骨幹端異形成症 (Shwachman-Bodian-Diamond症候群、SBDS) 回復性骨幹端異形成症1型 回復性骨幹端異形成症2型 骨幹端異形成症。Spahr型 骨幹端先端杯状異形成症(種々の型) 生殖器軟骨腫症(1型/2型) D-2-水酸化グルタール酸尿症を伴う骨幹端軟骨腫症	COL10A1 RMRP PTH1R [PTHR1] SBDS MMP13 MMP9	C		可能(応相談) 可能(応相談) 可能(応相談) 可能(応相談) 可能(応相談) 可能(応相談)
12. Spondylometaphyseal dysplasias (SMD) Odontochondrodysplasia (ODCD) Spondylometaphyseal dysplasia, Sutcliffe/corner fracture type SMD with severe genu valgum	12. 脊椎骨幹端異形成症(SMD) 歯牙軟骨形成不全症(ODCD) 脊椎骨幹端異形成症, Sutcliffe/corner fracture型 高度外反膝を伴うSMD				יייי איייין בייייי

ACP5

可能(応相談)

可能(応相談)

錐体・杆体ジストロフィを伴うSMD 網膜変性を伴うSMD, 脊柱型

脊椎軟骨内異形成症(SPENCD)

中間肢異形成症, Nievergelt型

中間肢異形成症, Kozlowski-Reardon型

SMD with retinal degeneration, axial type

Spondyloenchondrodysplasia (SPENCD)

Mesomelic dysplasia, Nievergelt type

Mesomelic dysplasia, Kozlowski-Reardon type

	Mesomelic dysplasia with acral synostoses (Verloes-David-Pfeiffer type) Mesomelic dysplasia, Savarirayan type (Triangular Tibia-Fibular Aplasia)	先端癒合症を伴う中間肢異形成症 (Verloes-David- Pfeiffer type) 中間肢異形成症, Savarirayan型 (三角形脛骨・腓骨無 形成症)				
	18. Bent bones dysplasias Campomelic dysplasia (CD) Stüve-Wiedemann dysplasia Kyphomelic dysplasia, several forms	18. 彎曲骨異形成症 屈曲肢異形成症 (CD) Stüve-Wiedemann骨異形成症 後彎肢異形成症, 各型	SOX9 LIFR	0	0 0	可能(応相談)
	19. Slender bone dysplasia Group 3-M syndrome (3M1) 3-M syndrome (3M2) Kenny-Caffey dysplasia type 1 Kenny-Caffey dysplasia type 2 Microcephalic osteodysplastic primordial dwarfism type 1/3 (MOPD1)	19. 狭細骨異形成症グループ 3-M症候群(3M1) 3-M症候群(3M2) Kenny-Caffey骨異形成症1型 Kenny-Caffey骨異形成症2型 小頭型骨異形成性原発小人症1型/3型(MOPD1)	CUL7 OBSL1 TBCE RNU4ATAC	0	0 0	
	Microcephalic osteodysplastic primordial dwarfism type 2 (MOPD2; Majewski type) Microcephalic osteodysplastic dysplasia, Saul-Wilson type IMAGE syndrome (Intrauterine Growth Retardation, Metaphyseal Dysplasia, Adrenal Hypoplasia, and Genital Anomalies) Osteocraniostenosis	小頭型骨異形成性原発小人症1型/3型(MOPD2; Majewski型) 小頭型骨異形成性異形成症, Saul-Wilson型 IMAGE症候群(子宮内胎児発育遅延, 骨幹端異形成, 副腎低形成, 性器異常) 骨頭蓋狭窄症	PONT		O(prenatalなし)	
	Hallermann-Streiff syndrome	育頭蓋狹乍症 Hallermann-Streiff症候群	GJA1			
— 196	20. Dysplasias with multiple joint dislocations Desbuquois dysplasia (with accessory ossification centre in digit 2) Desbuquois dysplasia with short metacarpals and elongated phalanges (Kim type) Desbuquois dysplasia (other variants with or without accessory ossification centre) Pseudodiastrophic dysplasia	20. 多発性脱臼を伴う骨異形成症 Desbuquois骨異形成症 (第2指の副骨化中心を伴う) 短い中手骨と長い指骨を伴うDesbuquois骨異形成症 (Kim 型) Desbuquois骨異形成症 (副骨化中心を伴う/伴わない 他の亜型) 偽性捻曲性骨異形成症	CANT1 CANT1		0	可能(応相談) 可能(応相談) 可能(応相談)
	21. Chondrodysplasia punctata (CDP) Group CDP, X-linked dominant, Conradi-Hünermann type (CDPX2)	21. 点状軟骨異形成症 (CDP) グループ CDP, X連鎖性優性遺伝型, Conradi-Hünermann型 (CDPX2)	EBP	0	0	
	CDP, X-linked recessive, brachytelephalangic type (CDPX1) CHILD (congenital hemidysplasia, ichthyosis, limb defects)	CDP, X連鎖性劣性遺伝型, 末梢骨短縮型 (CDPX1) CHILD症候群 (先天性片側異形成, 魚鱗癬様紅皮症, 四肢欠損)	NSDHL	0	0	可能(応相談)
	CHILD (congenital hemidysplasia, ichthyosis, limb defects) Greenberg dysplasia	CHILD症候群(先天性片側異形成, 魚鱗癬様紅皮症, 四肢欠損) Greenberg骨異形成症	EBP		0	
	Rhizomelic CDP type 1 Rhizomelic CDP type 2 Rhizomelic CDP type 3 CDP tibial-metacarpal type Astley-Kendall dysplasia	Greenberg有異形成症 近位肢型CDP 1型 近位肢型CDP 2型 近位肢型CDP 3型 CDP脛骨・中手骨型 Astley-Kendall骨異形成症	LBR PEX7 GNPAT AGPS	0	0 0 0 0 0	可能(応相談)
	22. Neonatal osteosclerotic dysplasias Blomstrand dysplasia Desmosterolosis Caffey disease (including infantile and attenuated forms) Caffey disease (severe variants with prenatal onset) Raine dysplasia (lethal and non-lethal forms)	新生児骨硬化性異形成症 Blomstrand骨異形成症 デスモステロール症 Caffey病 (乳児型・寛解型を含む) Caffey病 (出生前発症の重症型) Raine骨異形成症 (致死型および非致死型)	PTHR1 DHCR24 COL1A1 COL1A1 FAM20C		0 0 0 0 0	可能(応相談) 可能(応相談)
	23. Increased bone density group (without modification of bone shape) Osteopetrosis, severe neonatal or infantile forms (OPTB1)	骨変形を伴わない骨硬化性疾患グループ 大理石骨病、重症新生児型/乳児型(OPTB1)	TCIRG1		0	可能(応相談)
	Osteopetrosis, severe neonatal or infantile forms (OPTB4) Osteopetrosis, infantile form, with nervous system involvement (OPTB5)	大理石骨病,重症新生児型/乳児型 (OPTB4) 神経系罹患を伴う大理石骨病,乳児型, (OPTB5)	CLCN7 OSTM1	0	0	可能(応相談)

	Spondylo-ocular dysplasia Osteopenia with radiolucent lesions of the mandible Ehlers-Danlos syndrome, progeroid form Geroderma osteodysplasticum Cutis laxa, autosomal recessive form, type 2B (ARCL2B) Cutis laxa, autosomal recessive form, type 2A (ARCL2A) (Wrinkly skin syndrome) Singleton-Merten dysplasia	脊椎・眼異形成症 下顎骨X線透過性病変を示す骨減少症 Ehlers-Danlos症候群, 早老型 骨異形成性老人様皮膚症 皮膚弛緩症, 常染色体劣性遺伝型, 2B型 (ARCL2B) 皮膚弛緩症, 常染色体劣性遺伝型, 2A型 (ARCL2A), しわの多い皮膚症候群 Singleton-Merten症候群	B4GALT7 GORAB PYCR1 ATP6V0A2	0	0	
	26. Defective mineralization group Hypophosphatasia, perinatal lethal and infantile forms Hypophosphatasia, adult form Hypophosphatemic rickets, X-linked dominant Hypophosphatemic rickets, autosomal dominant Hypophosphatemic rickets, autosomal recessive, type 1 (ARHR1) Hypophosphatemic rickets, autosomal recessive, type 2 (ARHR2)	26. 骨石灰化障害を示すグループ 低フォスファターゼ症, 周産期致死型・乳児型 低フォスファターゼ症, 成人型 低リン血症性くる病, X連鎖性優性遺伝型 低リン血症性くる病, 常染色体優性遺伝型 低リン血症性くる病, 常染色体劣性遺伝型, 1型 (ARHR1) 低リン血症性くる病, 常染色体劣性遺伝型, 2型 (ARHR2)	FGF23 DMP1	0000	0 0 0 0 0	可能(応相談) 可能(応相談)
	Hypophosphatemic rickets with hypercalciuria, X-linked recessive Hypophosphatemic rickets with hypercalciuria, autosomal recessive (HHRH) Neonatal hyperparathyroidism, severe form Familial hypocalciuric hypercalcemia with transient neonatal	高Ca尿症を伴う低リン血症性くる病、X連鎖性劣性遺高Ca尿症を伴う低リン血症性くる病、常染色体劣性遺伝型(HHRH) 新生児上皮小体機能亢進症、重症型	CLCN5 DMP1 CASR		0 0 0	可能(応相談)
	hyperparathyroidism 27. Lysosomal Storage Diseases with Skeletal Involvement	一過性新生児上皮小体機能亢進症を伴う家族性高Ca 尿性高Ca血症	CASR		0	可能(応相談)
1	(Dysostosis Multiplex Group) Mucopolysaccharidosis type 1H / 1S Mucopolysaccharidosis type 2	27.骨変化を伴うリソソーム蓄積症(多発性異骨症グループ) ムコ多糖症 1H/1S型 ムコ多糖症 2型	IDUA IDS	0000	0	可能(応相談)
108 —	Mucopolysaccharidosis type 3A Mucopolysaccharidosis type 3B Mucopolysaccharidosis type 3C Mucopolysaccharidosis type 3D Mucopolysaccharidosis type 4A Mucopolysaccharidosis type 4B	ムコ多糖症 3A型 ムコ多糖症 3B型 ムコ多糖症 3C型 ムコ多糖症 3D型 ムコ多糖症 4A型 ムコ多糖症 4B型	HGSNAT GNS GALNS	000	0 0 0 0 0	
	Mucopolysaccharidosis type 6 Mucopolysaccharidosis type 7 Fucosidosis alpha-Mannosidosis	ムコタ糖症 6型 ムコ多糖症 7型 フコシドーシス アルファーマンノーシス	MAN2B1	00000	000000000000000000000000000000000000000	
	beta-Mannosidosis Aspartylglucosaminuria GMI Gangliosidosis, several forms Sialidosis, several forms	ベータ-マンノーシス アスパチルグルコサミン尿症 GMIガングリオドーシス、各型 シアリドーシス、各型 シアリドーシス、各型	MANBA AGA GLB1 NEU1	0	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	
	Sialic acid storage disease SIASD Galactosialidosis, several forms Multiple sulfatase deficiency Mucolipidosis II (I-cell disease), alpha/beta type	シアル酸蓄積症 SIASD ガラクトシアリドーシス, 各型 多種サルファターゼ欠損症 ムコ脂質症Ⅱ型 (I-cell病) ムコ脂質症Ⅲ型 (偽性Hurlerポリジストロフィー), アル	GNPTAB	0	0 0 0 0	
	Mucolipidosis III (Pseudo-Hurler polydystrophy), alpha/beta type Mucolipidosis III (Pseudo-Hurler polydystrophy), gamma type	ファ/ベータ型 ムコ脂質症Ⅲ型 (偽性Hurlerポリジストロフィー), ガンマ		0	0	
	28. Osteolysis Group Familial expansile osteolysis Mandibuloacral dysplasia type A Mandibuloacral dysplasia type B Progeria, Hutchinson-Gilford type Torg-Winchester syndrome Hajdu-Cheney syndrome	29. 骨溶解症グループ 家族性拡張性骨溶解症 下顎先端異形成症 A型 下顎先端異形成症 B型 早老症, Hutchinson-Gilford型 Torg-Winchester症候群 Hajdu-Cheney症候群	LMNA ZMPSTE24 LMNA MMP2 NOTCH2	000	O O O(prenatalなし)	
	Multicentric carpal-tarsal osteolysis with and without nephropathy	Bajdu-Gneney症候群 多中心性手根骨・足根骨溶解症(腎症を伴う/伴わな	INO I OFIZ		O	可能(応相談)