

ARTICLE

A novel intermediate mucopolipidosis II/III $\alpha\beta$ caused by GNPTAB mutation in the cytosolic N-terminal domain

Jules G Leroy¹, David Sillence², Tim Wood¹, Jarrod Barnes³, Robert Roger Lebel⁴, Michael J Friez¹, Roger E Stevenson¹, Richard Steet³ and Sara S Cathey^{*,1}

Mucopolipidosis (ML) II and ML III $\alpha\beta$ are allelic autosomal recessive metabolic disorders due to mutations in *GNPTAB*. The gene encodes the enzyme UDP-GlcNAc-1-phosphotransferase (GNPT), which is critical to proper trafficking of lysosomal acid hydrolases. The ML phenotypic spectrum is dichotomous. Criteria set for defining ML II and ML III $\alpha\beta$ are inclusive for all but the few patients with phenotypes that span the archetypes. Clinical and biochemical findings of the 'intermediate' ML in eight patients with the c.10A>C missense mutation in *GNPTAB* are presented to define this intermediate ML and provide a broader insight into ML pathogenesis. Extensive clinical information, including radiographic examinations at various ages, was obtained from a detailed study of all patients. *GNPTAB* was sequenced in probands and parents. GNPT activity was measured and cathepsin D sorting assays were performed in fibroblasts. Intermediate ML patients who share the c.10A>C/p.K4Q mutation in *GNPTAB* demonstrate a distinct, consistent phenotype similar to ML II in physical and radiographic features and to ML III $\alpha\beta$ in psychomotor development and life expectancy. GNPT activity is reduced to 7–12% but the majority of newly synthesized cathepsin D remains intracellular. The *GNPTAB* c.10A>C/p.K4Q missense allele results in an intermediate ML II/III with distinct clinical and biochemical characteristics. This delineation strengthens the utility of the discontinuous genotype–phenotype correlation in ML II and ML III $\alpha\beta$ and prompts additional studies on the tissue-specific pathogenesis in GNPT-deficient ML.

European Journal of Human Genetics (2013) 0, 000–000. doi:10.1038/ejhg.2013.207

Keywords: mucopolipidosis; phosphotransferase; storage disease; *GNPTAB*

INTRODUCTION

Mucopolipidosis II (ML II; I-cell disease; MIM 252500) and mucopolipidosis III $\alpha\beta$ (ML III; pseudo-Hurler-polydystrophy; MIM 252600) are allelic autosomal recessive metabolic disorders directly correlated with homozygous or compound heterozygous mutant *GNPTAB* genotypes. The *GNPTAB* gene encodes the α and β polypeptides in the hexameric enzyme UDP-GlcNAc-1-phosphotransferase (GNPT; IUBMB no. 2.7.8.17). The γ polypeptide is encoded by the *GNPTG* gene. The enzyme complex catalyzes the first step in the biosynthesis of the mannose-6-phosphate (M6P) recognition marker in the oligomannosyl-type N-linked glycans in lysosomal acid hydrolases. Without the M6P marker most soluble hydrolases are not targeted to the lysosomal compartment in connective tissues, resulting in impaired lysosomal degradation.

GNPTAB genotypes, either homozygous or compound heterozygous for mutant alleles with 'amorph' or 'null' effect (frameshift and nonsense mutations), result in the early-onset, severe ML II. The later-onset ML III $\alpha\beta$ with prolonged course is the morbid result in patients with at least one 'hypomorph effect' allele (missense and most splice mutations) in the *GNPTAB* mutant genotype. The phenotypic spectrum of patients with GNPT-deficiency disorders is dichotomous rather than continuously variable. ML II and ML III $\alpha\beta$ are phenotypically distinguishable. Within the largest published cohort of ML patients, the phenotype of only a few subjects straddled the criteria set for defining ML II and ML III $\alpha\beta$.¹

The purpose of this report is to present and discuss the clinical and biochemical findings of the 'intermediate' ML in eight patients with the c.10A>C missense mutation in exon 1 in their compound heterozygous *GNPTAB* genotype. The amino acid lysine is replaced with glutamine, p.K4Q (designated K4Q hereafter). The name 'Webb type ML' is proposed following the explicit permission for use of the surname of one of the families and honoring all participating families and patients in this study. The findings corroborate the overall genotype–phenotype correlation, underscore the predictive power of mutation screening and illustrate the potential for generating heuristic hypotheses on the pathogenesis of ML.

SUBJECTS AND METHODS

Probands 1, 2 and 3 are isolated patients in three unrelated families. Only patient 3, who died at the age of 23 6/12 years, was never examined by at least one of the authors. However, the wealth of clinical data made available by the mother prompted inclusion in the core group of subjects reported earlier and in this study.¹ The postmortem study of patient 3 was recently published as the report of autopsy findings in a patient with ML III $\alpha\beta$.² Patients 4–8 are siblings. The earlier part of the disorder's course in the older siblings was reported by Kozłowski *et al.*³ All sibs were included in the recent report by David-Vizcarra *et al.*⁴ who recognized the affected as examples of ML intermediate between ML II and ML III $\alpha\beta$. The parents in each of the four families were non-consanguineous.

In addition to plasma samples and fibroblast cell lines representing the eight subjects, cultured fibroblasts from two additional patients, one homozygous

¹Department of Clinical Genetics, Greenwood Genetic Center, Charleston Office, North Charleston, SC, USA; ²The Children's Hospital at Westmead, Westmead, New South Wales, Australia; ³Complex Carbohydrate Research Center, University of Georgia, Athens, GA, USA and ⁴State University of New York Upstate Medical School, Syracuse, New York, USA

*Correspondence: Dr SS Cathey, Department of Clinical Genetics, Greenwood Genetic Center, Charleston Office, 3520 W. Montague Avenue, Suite 104, North Charleston, SC 29418, USA. Tel: +1 843 746 1001; Fax: +1 843 746 1002; E-mail: scathe@ggc.org

Received 31 January 2013; revised 5 July 2013; accepted 11 July 2013

and one compound heterozygous for the K4Q missense mutation, were obtained from NIGMS Human Genetic Cell Repository at Coriell Institute (Camden, NJ, USA) and included in the biochemical studies. No clinical information except the diagnosis of ML III was available. The mutant genotypes in patients and cell lines are presented in Table 1.

The activity of selected lysosomal acid hydrolases was assayed in plasma and cultured fibroblasts according to the methods described by Thomas *et al.*⁵ Metabolic labeling and sorting of the lysosomal hydrolase cathepsin D in control and patient fibroblasts were performed as described previously.^{6,7} Briefly, cell monolayers at about 60–80% confluency were washed twice with phosphate-buffered saline and methionine starved using cysteine/methionine-free Dulbecco's modified Eagle's medium for 30 min, followed by a pulse label for 1 h with 1 ml of 1 mCi/ml Tran³⁵S-label (MP Biomedicals) in the same media. Excess unlabeled methionine (10 mM final concentration) was added to initiate a 4-h chase. The amount of cathepsin D sorted was determined by overnight immunoprecipitation of equivalent aliquots of the lysed cell and media high-speed supernatants at 4 °C with rabbit anti-human cathepsin D antiserum and protein A-Sepharose beads. After five washes with buffer containing 1 M KCl and 1% Triton X-100, immunoprecipitates were subjected to 10% SDS-PAGE under nonreducing conditions. Gels were dried and exposed on an autoradiography film. Quantification of the various bands was performed using Image J software. The percentage of cathepsin D that remains intracellular was calculated as the densitometric ratio of the intracellular forms (both intermediate and mature) to the sum of the intracellular and secreted (precursor) forms.

GlcNAc-1-phosphotransferase activity was measured in microsomal preparations from fibroblasts as described earlier.^{8,9} Briefly, cells were lysed by sonication in Tris buffer pH 7.5 containing 0.25 M sucrose and microsomes prepared by ultracentrifugation. Reactions using 10 mM α -methylmannoside as an acceptor and 3H-UDP-GlcNAc (along with 200 μ M 'cold' UDP-GlcNAc) as substrate were conducted in the following reaction buffer: 50 mM Tris pH 7.5, 0.3% Triton X-100, 50 mM *N*-acetylglucosamine, 1 mM DTT, 20 mg/ml bovine serum albumin, 10 mM sodium molybdate, 1 mM MgCl₂, and 1 mM MnCl₂. Activity units are calculated as pmoles of 3H GlcNAc transferred per hour. Protein concentration was measured using the micro BCA protein assay kit (Pierce, Rockford, IL, USA).

Screening for the *GNPTAB* mutations by PCR and sequencing protocols was performed according to the methods reported earlier.¹

RESULTS

Clinical phenotype

Patient 1. Patient 1 was delivered at term after an uncomplicated pregnancy (panels A1 and A2 in Figure 1). Size was appropriate for gestational age (Table 2). Early features included plagiocephaly treated with helmeting, kyphosis noted at approximately 9 months and regurgitant cardiac valves confirmed by echocardiogram at 12 months. Skeletal survey was performed at 20 months when a decreased range of motion in the shoulders prohibited lifting the

arms above the head. The radiographs showed features of dysostosis multiplex (DM) that included 33° kyphosis, varus angulation of the proximal humeri, bowed, dysplastic radii and ulnae, proximal and distal tapering of the widened metacarpals, and marked coxa valga and subluxation of the hips. The diagnosis of ML II was assigned at 22 months. While anthropometric measurements were normal in the first year of life, growth failure was apparent from his second year. At 1 year, length was 76 cm (50th percentile). The maximal height of 80.5 cm (mean for a 16-month-old male) was recorded at 7.5 years. Speech and expressive language development were normal but unaided walking was not achieved until 20 months. Cognitive development was delayed. Formal assessment at 12 11/12 years showed a global intellectual ability score of 39, comparable to an age equivalent of 5 5/12 years. Obstructive sleep apnea was treated with continuous positive airway pressure at night, beginning before 5 years. Recurrent infections led to one hospitalization for pneumonia and three sets of myringotomy tubes by 8 years. Several surgical attempts were made to stabilize his progressive thoracolumbar kyphosis, with operations at 3, 6, and 10 years. Surgery for relief of bilateral carpal tunnel syndrome was performed at 7 years. Post-operative extubation was consistently difficult. Prolonged ventilator support with slow weaning was required. When examined at 12 7/12 years progressive joint contractures had decreased measured height to 78.7 cm (mean for a 15-month-old male), weight was 15 kg (mean for a 3-year-old male), and head circumference 51.5 cm (mean for a 6-year-old male). He wore glasses for nearsightedness. Dysmorphic features included flat and coarse facies, ridged palate, thickened gums, broad and short hands held in ulnar deviation, broad and camptomelic fingers. There was a small umbilical hernia and diastasis of the straight abdominal muscles without organomegaly. Despite joint restriction, the boy was mobile, interactive and actively playful. He conversed appropriately with hoarse voice.

Patient 2. Patient 2 was delivered by cesarean section for transverse lie following a pregnancy complicated by hyperemesis in the first trimester (panel B in Figure 1). Ultrasound examination at 18 weeks gestation was normal. Neonatal anthropometrics were normal (Table 2). Features noticed on the first day of life included unusual facies with micrognathia, contractures at the knees and hips but hypermobile wrists, slender fingers and narrow feet. Slow weight gain in infancy was improved temporarily when breastfeeding was supplemented with bottle-feeding. Impaired growth was apparent by 22 months. Speech developed normally. He had significant gross motor delay, persistent truncal hypotonia and mild thoracolumbar kyphoscoliosis noticed at 12 months. He did not sit unassisted until 18 months and walked at 30 months. At 33 months, he did not have organomegaly, corneal clouding or gingival swelling, but lysosomal storage disease was suspected based on his coarse facies, short stature, and skeletal changes. Testing revealed increased activity of multiple lysosomal enzymes in plasma. The initial diagnosis of ML II was assigned, but by age 40 months the overall clinical phenotype was more in support of intermediate ML and called 'type II/III'. Examination at 10 2/12 years showed an alert and conversant, ambulatory boy with stature 86.5 cm (mean for a 23-month-old male), weight 16.4 kg (mean for a 12-month-old male), and head circumference 51.5 cm (mean for a 6-year-old male). Relevant findings included normocephaly, coarse facies, gingival hypertrophy, narrow thorax, soft ejection murmur, mildly prominent abdomen without organomegaly, umbilical hernia, thoracolumbar kyphoscoliosis, severely limited range of motion in all joints, short claw-like hands in ulnar deviation, and broad camptomelic fingers.

Table 1 *GNPTAB* mutant genotype in *GNPTAB* c.10A>C/p.K4Q ML patients and in mutant fibroblast lines

Patient	Paternal allele	Maternal allele
1	Exon 1: c.10A>C/p.K4Q	Exon 13: c.1964delC
2	Exon 11: c.1399delG	Exon 1: c.10A>C/p.K4Q
3	Exon 13: c.2190delT	Exon 1: c.10A>C/p.K4Q
4–8	Exon 1: c.10A>C/p.K4Q	Exon 11: c.1399delG
<i>Line</i> ^a		
GM00113	Exon 1: c.10A>C/p.K4Q	Exon 1: c.10A>C/p.K4Q
GM01586	Exon 19: 3665_3666delTC	Exon 19: 3665_3666delTC

^aAdopted from Human Genetic Cell Repository, Coriell Institute for Medical Research, Camden, NJ, USA; genotypes.



Figure 1 *GNPTAB* c.10A>C/p.K4Q ML clinical features in three unrelated patients: short stature, normocephaly, coarse facies, limited mobility in all joints, short and broad hands with fingers postured in ulnar deviation. A₁ and A₂: Patient 1 at 8 and 12 years; B: Patient 2 at 6 years; C₁ and C₂: Patient 3 at 8 and 20 years.

Q6 *Patient 3.* Patient 3 was the third of three sons born to a Japanese father and Caucasian mother (panels C₁ and C₂ in Figure 1). Pregnancy was uncomplicated until 4 weeks prior to the expected delivery date when rupture of membranes and labor led to vacuum-assisted vaginal delivery. Size was appropriate for gestational age (Table 2). By 9 months, parents regarded his development to be slower than that in his healthy siblings and noted differing physical features, including proptotic eyes, short neck, relatively large head, narrow shoulders, and protruding abdomen. Speech was late (first words at 15 months) and difficult to understand. He achieved unaided walking at 18 months despite stiff hips and knees. By 23 months, he had limited range of motion at the shoulders, wrists and fingers, and thoracolumbar kyphoscoliosis. ML II was first suspected at 26 months; ML III was his diagnosis at 29 months. Minimal statural growth occurred after age 3 years. He had speech therapy and was in special education classes throughout the years of schooling. Testing of academic achievement at 14 years revealed a school performance at the third to fourth grade level. At 11 years, he had mild thickening of aortic and mitral valves, mitral regurgitation and left ventricular hypertrophy. Cardiac and pulmonary function

worsened in the second decade. At 21 years, the left ventricular ejection fraction was below 60% and associated with mild to moderate regurgitation at all cardiac valves. He had severe sleep apnea by mid-teens and required continuous supplemental oxygen by 22 years. He died at 23 years. Autopsy confirmed severe dilated hypertrophic cardiomyopathy.²

Patients 4–8: common features in sibling patients. Five of seven children born to healthy parents were affected (Figure 2; Table 2). The two surviving affected siblings (patients 5 and 8) are ages 28 and 15 years. Patients 4, 6, and 7 died at 21, 24, and 14 years, respectively. Placental abruption at term complicated one pregnancy. The third affected child, a male, was diagnosed prenatally and had an unaffected female twin. No ultrasound abnormalities had been detected in either fetus. Kyphosis and dislocated hips were diagnosed in the firstborn child (patient 4) when she was 9 months old. Dysmorphic facial features and joint contractures were subsequently recognized in infancy in each of the affected siblings. The diagnosis of ML II was made simultaneously in the two oldest children at 26 months and 7 months. All of the children were of appropriate size for gestational

Table 2 Clinical features in eight patients with *GNPTAB* c.10A>C/p.K4Q ML

Patient	1	2	3	4	5	6	7	8
Gender	Male	Male	Male	Female	Female	Male	Female	Female
Pregnancy term (weeks)	39	39	36	39	41	40	40	42+
Birth weight (g)	3200	3430	2390	3060	3290	2520	3290	3970
Birth length (cm)	53	53	49	54	54			
Birth OFC (cm)	33	34		35	34			
<i>Age at first signs</i>								
Craniofacial	1 day	1 day	9 months	1 day	1 day	1 day	1 day	1 day
Skeletal	6 months	1 day	9 months	8 months	5 months	1 day	1 day	1 day
Age at ML diagnosis	22 months	33 months	23 months	26 months	7 months	1 month	2 weeks	2 weeks
<i>Age at motor milestones</i>								
Sitting	12 months	18 months	12 months	18 months	9 months	9 months	9 months	9 months
Crawling	14 months	18 months		36 months	12 months	11 months	12 months	12 months
Walking	20 months	30 months	18 months	7 years	21 months	42 months	42 months	42 months
Max height (cm)	80.5	87.0	88.0		90		84.5	87.8
Age at max height (years)	7.5	12	22		26		13	15
Age at death			23 years	21 years		24 years	14 years	

Patients are from four unrelated families. Patients 4–8 are siblings.



Figure 2 *GNPTAB* c.10A>C/p.K4Q ML clinical features in five siblings: Similar findings as in Figure 1 are shown early and late in disorder's natural course. D: Patient 4 at 21.5 years; E: Patient 5 at 20 years; F: Patient 6 at 17.5 years; G: Patient 7 at 7.5 years; H: Patient 8 at 6 years.

age, but statural growth slowed from infancy and ceased in early childhood. Receptive and expressive language development occurred at a near normal pace but neuromotor development was delayed. Ambulation was not achieved until age 7 years in patient 4, who had severe congenital flexion contractures at the knees. Patient 7 maintained independent ambulation only briefly. Cognitive development was impaired, and all participated in special education schooling. None truly mastered reading but could copy letters and words. Recurrent bouts of otitis contributed to conductive hearing loss and

led to repeated sets of myringotomy tubes in all. All patients had surgery for carpal tunnel syndrome. Systolic heart murmurs were present from early childhood. Echocardiography revealed slow thickening and gradually worsening incompetence of the mitral and aortic valves. Later in the clinical course, enlargement of the left ventricular wall and dilatation of the left atrium and ventricle were noticed. From infancy, breathing was louder, more superficial and faster than normal. Adenoidectomy was performed in some. Sleep apnea, obstructive and central, was treated with continuous positive

airway pressure. There was limited development of secondary sexual characteristics; only one of the females ever had menses. Disease progression was characterized by severe bone and joint pain even when at rest. Periodic intravenous infusion of biphosphonate provided transient pain relief. All patients developed radiographic and neurologic signs of narrowing of the spinal canal. Although disease progression was apparent in every system, the morbid consequences of cord compression dominated end-of-life issues.

The clinical diagnosis of intermediate mucopolidosis (ML II/III) was reached in the two prior reports that included clinical and radiological data from affected members of this family.^{3,4}

The radiographic phenotype

The osteochondrodysplasia (OCD) documented radiographically in this group of ML patients fits within the concept of DM defined and illustrated by Spranger *et al.*¹⁰ The skeletal radiographs in this group of patients at various ages show the remarkably consistent DM that is congruent with ML II and significantly different from the ML III α/β radiologic phenotype (Table 3; Figures 3 and 4).^{1,3,4,10,11}

In the lateral skull films available from late childhood only the sella turcica has an elongated shape, whereas in age-matched and older ML III α/β patients it maintains the normal configuration. The thoracolumbar vertebrae demonstrate a much foreshortened and near ovoid shape with convex upper and lower borders, and concave anterior and posterior irregular rims. At least one vertebra, either T12 and/or L1 at the top of the kyphotic spine deformation appears more hypoplastic,

insufficiently ossified and often triangular (Figure 3). The latter feature is consistently present in ML II, but absent in some ML III α/β patients. In the anteroposterior thoracic films, the ribs are considerably widened in the lateral and anterior segments, but abnormally narrow near the vertebral insertion (not shown). The pelvis is most obviously osteopenic with severely slanted acetabular roofs, narrow basilar portions of the ilia and relatively large and elongated ischial and pubic bones, features consistently present in ML II and also ML III α/β . All long bones are short and poorly tubulated with thick cortices and small, coarsely trabeculated epiphyses. The knee epiphyses are small and almost rectangular. Ossification of the lateral portion of the tibial submetaphyseal zone is overmodeled. Changes in the hands and wrists (Figure 4) fit into the severe side of the DM spectrum. All epiphyses in the tubular hand bones and the carpal bones remain small and irregular. The metacarpal diaphyses are excessively widened and ends are constricted. While nearly congruent with the changes in ML II, the hand skeleton in these intermediate ML patients differs significantly from the much milder OCD in ML III α/β hand bones (Figure 4). In the longest surviving patients, the radiographic findings are altered by the worsening osteopenia.

Biochemical and molecular results

Acid glycosidase levels in serum and cells. The activity of total β -D-hexosaminidase, α -L-fucosidase and β -glucuronidase was significantly increased in plasma and decreased in cultured fibroblasts derived from patients. These results overlap with the ranges of hydrolase activity observed in ML II and ML III α/β . These findings do not contribute to distinguishing the *GNPTAB* c.10A>C/p.K4Q ML patients from either of the archetype GNPT-deficient disorders.^{1,12,13}

Assay of UDP-GlcNAc 1-phosphotransferase (GNPT). The results of the GNPT assay are presented in Table 4. The activity of the phosphotransferase towards the substrate α -methylmannoside is reduced to a range of 7–12% in the *GNPTAB* c.10A>C/p.K4Q ML patients, about half of that in the K4Q homozygous patient fibroblasts (GM00113), which is to be expected.¹⁴ The significantly higher GNPT activity in the K4Q samples compared with classic ML II supports the distinct nature of the *GNPTAB* c.10A>C/p.K4Q ML.

Sorting efficiency of cathepsin D. The phenomenon of missorting of immature lysosomal enzymes and their appearance outside the cell was studied in nascent cathepsin D by pulse-chase immunoprecipitation experiments. Cathepsin D is a highly mannose phosphorylated lysosomal protease that is hypersecreted from the cell when mannose phosphorylation is impaired. The pulse/chase assays thus provide an opportunity to gauge the amount of newly synthesized cathepsin D that remains intracellular *versus* that secreted into the media. Whereas the level of intracellular cathepsin D in fibroblasts from either controls or obligate heterozygous parents exceeds the normal 90% (Figure 5), ML II-type 'I-cells' (GM01586) retain only about 20% of newly synthesized cathepsin D. The fibroblasts from our patient 2 and the homozygous K4Q cells (GM0113) exhibit surprising levels of intracellular cathepsin D (81% and 65%, respectively). The comparatively reduced percentage in the GM00113 cells is attributed to the late passage number and consequent slow-growing nature of this culture. Greater heterogeneity in all forms of cathepsin D was detected in patient 2 and GM00113, which may reflect altered trafficking and/or processing of this hydrolase within the secretory pathway.

GNPTAB mutation screening. The *GNPTAB* mutation analysis data in patients 1–3 and 8 were previously reported.¹ Table 1 also lists the

Table 3 Distinctive phenotypic findings in *GNPTAB* c.10A>C/p.K4Q ML

Finding	ML II	K4Q ML	ML III
<i>Clinical</i>			
Coarsening of facies	++	+	±
Impaired growth	++	++	+
Short hands, fingers	++	++	±
Limited motion range in joints (early childhood)	++	+	±
Unaided ambulation	– ^a	+ ^b	+ ^b
Speech (normal, near normal)	–	+	+
Intellectual disability	++	+	±
Survival beyond childhood	–	+	+
<i>Radiographic</i>			
Overall dysostosis multiplex before age 3 years (skeletal survey)	+++	+++	+
Sella turcica elongated (lateral skull)	^c	+	–
Thoracolumbar vertebrae foreshortened (lateral spine)	++	++	–
Deformation of carpal, metacarpal, phalangeal bones (hands, wrists)	++	++	±
<i>Transient phenomena in the perinatal, infantile periods</i>			
Periosteal 'cloaking' alongside diaphyses of large long bones	++	^d	–
Periarticular punctate calcifications (prenatal ultrasound)	++ ^e	^d	–

^aIndependent walking rarely and very transiently achieved.

^bIndependent ambulation consistently lost from adolescence because of hip dysplasia and painful osteopenic dystrophy.

^cShort life span may preclude elongation of sella turcica in ML II.

^dDocuments for proof or exclusion not available.

^eSonographic finding in fetal ML II; inconsistent in ML II infants.

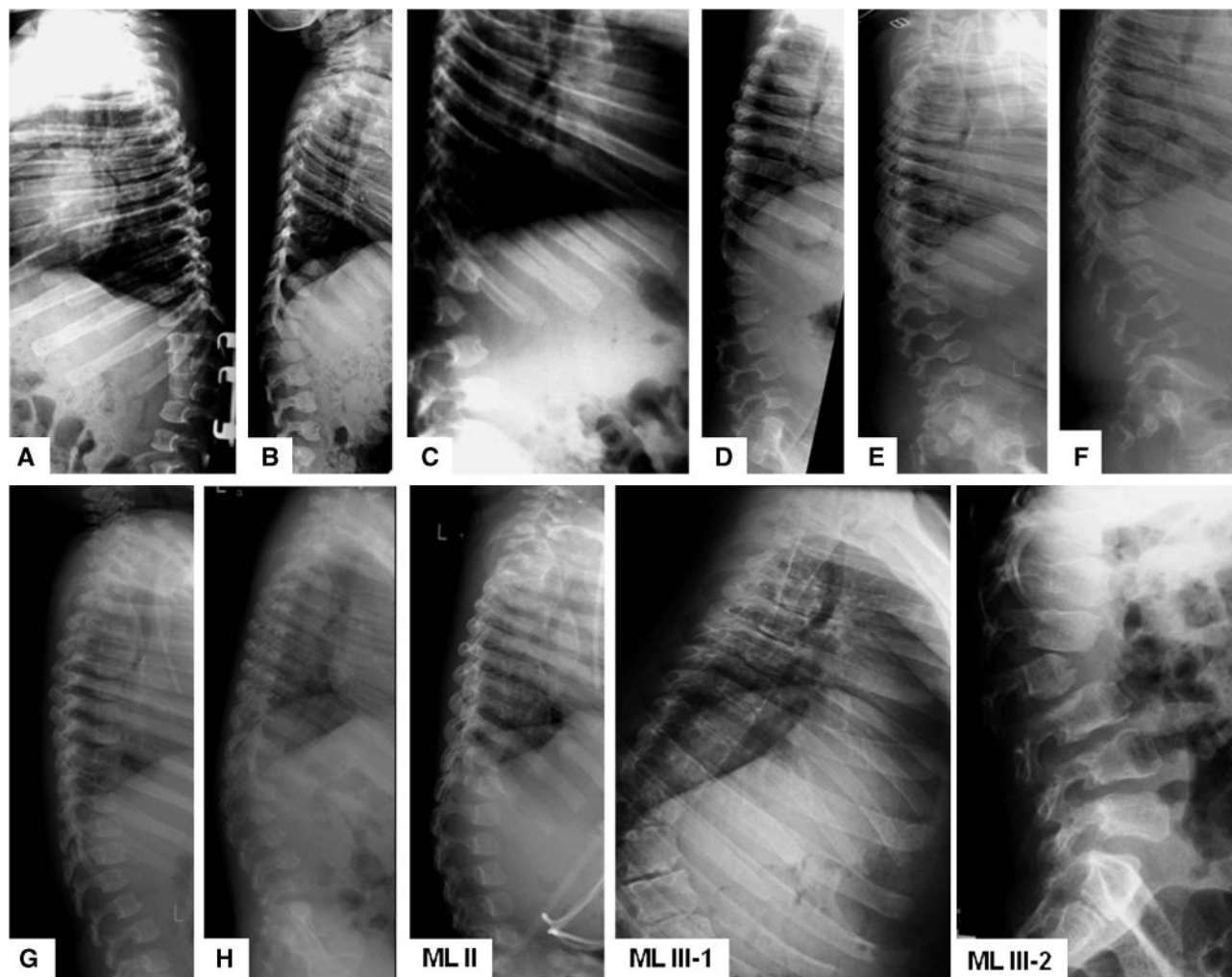


Figure 3 Lateral radiographs of spinal columns in eight patients with *GNPTAB* c.10A>C/p.K4Q ML compared with one ML II and two ML III α/β patients. Thoracolumbar vertebral bodies remain significantly foreshortened with concave frontal and convex upper/lower margins; delayed and deficient ossification. Similar findings are seen in ML II. In ML III α/β vertebral bodies are oblong, and show mild platyspondyly and irregular margins higher anteriorly than posteriorly. In all types of ML ossification is most deficient in either T12 or L1 or in both, often at the top of thoracolumbar kyphoscoliosis. A: Patient 1 at 8 years; B: Patient 2 at 6 years; C: Patient 3 at 9 years; D: Patient 4 at 19 years; E: Patient 5 at 19 years; F: Patient 6 at 18 years; G: Patient 7 at 7 years; H: Patient 8 at 4 years; ML II: female ML II patient at 4 years; ML III-1: male with ML III α/β at 18 years; ML III-2: female with ML III α/β at 13 years.

mutant *GNPTAB* genotypes for the two fibroblast lines obtained from the Coriell Institute.

DISCUSSION

Of the 61 probands included in the report by Cathey *et al*,¹ the clinical features in only a few patients with *GNPT*-deficient ML straddled the boundaries of the variability spectra set for delineation of ML II and of ML III α/β .¹ Upon referral for *GNPTAB* mutation screening, all had been diagnosed as ‘intermediate ML’ or ‘ML II/III’. The majority of these ‘intermediate’ patients was found to have a compound heterozygous mutant genotype that consistently included c.10A>C/p.K4Q missense mutation in exon 1 and a frameshift mutation in a downstream exon (Table 1). This group of ML patients has a distinct and remarkably consistent phenotype that shares physical (Figures 1 and 2) and radiographic (Figures 3 and 4) features with ML II, but more closely resembles the ML III α/β phenotype regarding life expectancy, psychomotor development and degree of intellectual disability.

The *GNPTAB* missense (MS) mutation c.10A>C/p.K4Q causes a unique intermediate type ML. The two cytosolic domains of *GNPTAB* at the N- and C-termini contain motifs that mediate the export or trafficking of the enzyme through the endoplasmic reticulum and Golgi. The importance of the N-terminal domain has been highlighted by recent work from Franke *et al*¹⁵ showing that the lysine residue at position 4—the same residue mutated in the patients described in this report—affects a neighboring dileucine sequence responsible for the export of the GlcNAc-1-phosphotransferase $\alpha\beta$ precursor from the endoplasmic reticulum.¹⁵ The possibility that other mutations may be found to result in an identical phenotype cannot be excluded. It remains to be seen whether other mutations within the N-terminal domain, or perhaps the C-terminal cytosolic domain, are associated with the unique clinical and biochemical features described in the patients reported here. This distinct ML subtype may be associated with other mutations that directly impact the cytosolic sorting motifs, a unique alteration of *GNPTAB*. Differences within the organization of the secretory pathway and

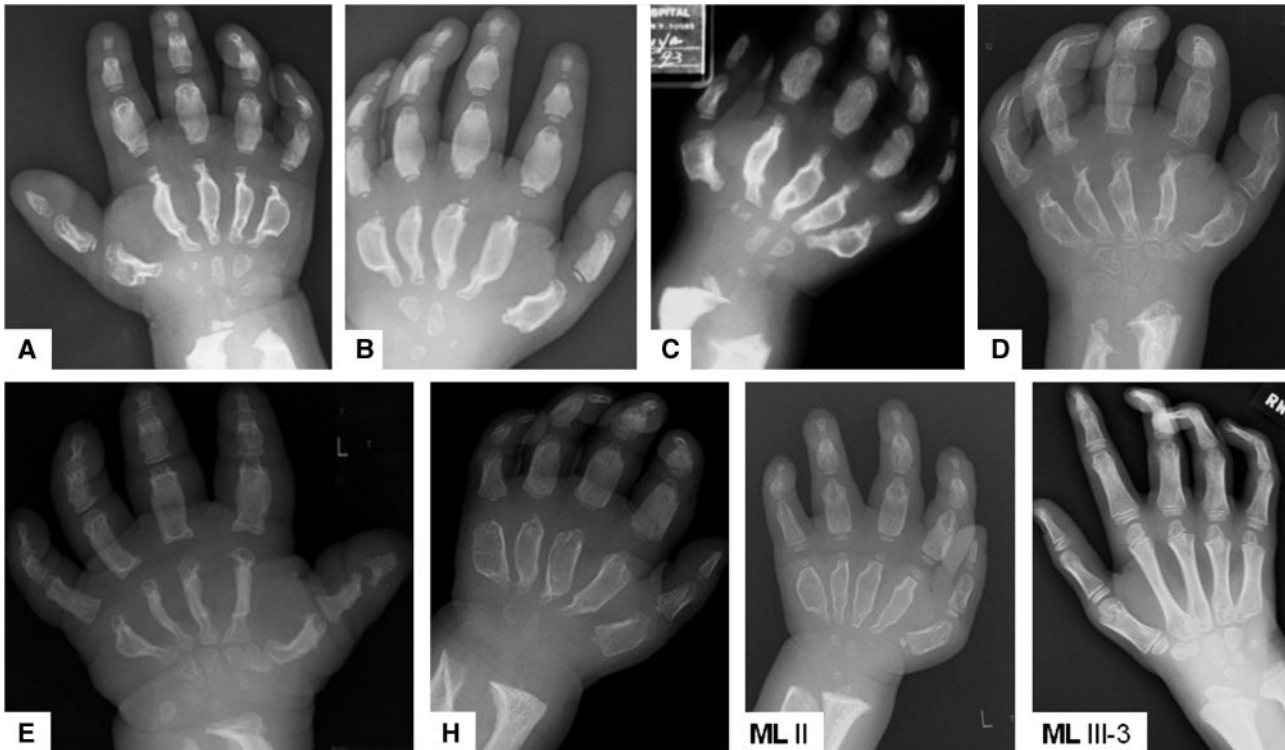


Figure 4 Hand radiographs in *GNPTAB* c.10A>C/p.K4Q ML patients: comparison with ML II and ML III $\alpha\beta$. DM in the hands is similar to that in ML II and distinctly more severe than in ML III $\alpha\beta$. Figures show shortening and diaphyseal widening in all tubular hand bones; severely delayed and deficient ossification of epiphyses, carpal bones and the distal ulnar and radial epiphyses; and apparent regression of mid-diaphyseal widening in older patients concomitant with progressive generalized osteopenia. A: Patient 1 at 8 years; B: Patient 2 at 4 years; C: Patient 3 at 9 years; D: Patient 4 at 9 years; E: Patient 5 at 19 years; H: Patient 8 at 6 years; ML II: female with ML II at 4 years; ML III-2: female with ML III $\alpha\beta$ at 13 years.

Table 4 GNPT activity for *GNPTAB* c.10A>C/p.K4Q ML patients and other patient fibroblast lines

Patient/ cell line ID	GNPT activity (pmol/mg/h)	% of average control activity	Genotype	Clinical diagnosis
Patient 2	1.79	11.1	p.K4Q/c.1399delG	Intermediate
Patient 5	1.86	11.6	p.K4Q/c.1399delG	Intermediate
Patient 8	1.14	7.1	p.K4Q/c.1399delG	Intermediate
GMO0113	3.14	19.5	p.K4Q/p.K4Q	'ML III' ^a
ML III	2.13	13.2	p.S399F/ c.3335+6T>G	ML III
affected GMO1586	0.31	1.9	c.3665_3666delTC/ c.3665_3666delTC	ML II
Control 1	13.8	—	—	Control
Control 2	18.5	—	—	Control
Control 3	16.0	—	—	Control

The three control activities were averaged and used to compare the relative activity in patient samples. The GNPT activity for GMO1586 and Patient 2 represents the average of two independent experiments. All other activities were single determinations.
^aThe GNPT activity result for GMO0113 is in good agreement with earlier data from two groups who analyzed these cells.^{16,17} Unfortunately, no clinical information could be obtained for this patient.

the trafficking machinery within certain cell types may lead to differences in GlcNAc-1-phosphotransferase trafficking. We believe such differences may help explain the tissue-specific phenotypes seen within the *GNPTAB* c.10A>C/p.K4Q ML patients.

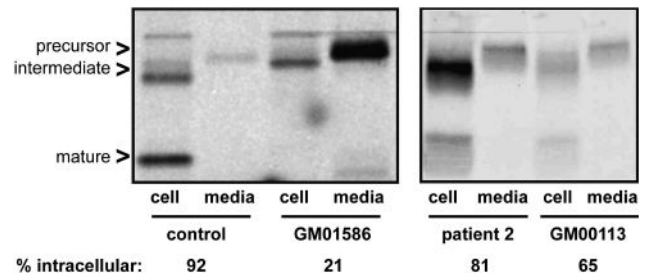


Figure 5 Sorting of newly synthesized cathepsin D in control and ML fibroblasts GMO1586 (ML II), patient 2 (K4Q compound heterozygote), and GMO0113 (K4Q homozygote). Autoradiographs of ³⁵S-labeled immunoprecipitates of cell and secreted cathepsin D are shown. The percentage of intracellular enzyme is determined by dividing the amount of cell-associated cathepsin D by the total cathepsin D (cell-associated and secreted).

The reduced intracellular and much enhanced plasma activity of several acid glycosidases in this cohort does not differ significantly from the corresponding data obtained in the reference phenotypes, ML II and ML III $\alpha\beta$. The GNPT assays (Table 4) confirm earlier reports in leukocytes and fibroblasts that activity of the causative enzyme is nearly completely absent in ML II and between 2 and 20% of normal in ML III $\alpha\beta$.^{1,8,9,16} The residual GNPT activity in each of the K4Q compound heterozygous ML lines is higher than that in ML II-derived fibroblasts and almost matches that in the ML III $\alpha\beta$ line. Remarkably, but not unexpectedly, GNPT activity in the homozygous

K4Q mutant cell line is 19.5% of that in controls and confirms the results previously obtained by others.¹⁷ This observation proves that the K4Q *GNPTAB* allele is indeed pathogenic but fairly leaky at least in cultured fibroblasts. The molecular characterization and trafficking of the enzyme may be further elucidated by additional studies of the mechanism by which this mutation alters GNPT activity.

The subnormal yet still effective maintenance of intracellular cathepsin D levels demonstrates the utility of this assay in distinguishing this specific intermediate type ML at least from the ML II reference phenotype. The fact that the majority of newly synthesized cathepsin D remains inside the cell in the *GNPTAB* c.10A>C/p.K4Q ML patients is surprising, and inconsistent with both the severity of the phenotype and the elevated plasma levels of lysosomal enzymes. However, these patient cells do retain substantial activity towards the simple sugar substrate, α -methylmannoside. We cannot rule the possibility that this mutation results in hydrolase-specific effects on lysosomal sorting. Further research on cathepsin-sorting efficiency may provide insight into the early failure of statural growth, nearly as severe as in ML II. The robust lysosomal enzyme missorting observed in ML II cells may extend to other cathepsins (for example, cathepsin K) known to play central roles in skeletal development. The relatively high efficiency of cathepsin D sorting may explain the moderate cognitive disability associated with this genotype. Roles for cathepsin D have been suggested in neural development and degeneration, and maintenance of near-normal intracellular levels of this protease may confer a protective effect within the brain of these patients.^{18–20}

Delineation of this c.10A>C/p.K4Q GNPT-dependent intermediate ML has more than merely clinical and prognostic significance. The partial ‘uncoupling’ of physical and cognitive disability represents the apparently rare exception to the rule that nearly complete absence of GNPT activity yields the ML II phenotype and more residual activity results in ML III $\alpha\beta$.¹ The severe DM, including the transient skeletal changes in infancy, is compatible with the ML II archetype. However, the specific intermediate type ML we describe is associated with GNPT activity and extension of life expectancy compatible with the MLIII $\alpha\beta$ archetype, and thus illustrates the progressively morbid clinical course not granted to the ML II patient. Manifestations include progressive and painful osteopenia, areas of frank osteolysis, and severe sclerosis in fasciae with contractures in tendons and other periarticular structures. Progressive destruction of the weight-bearing skeleton occurs both in the lower third of the ilia and particularly the femoral necks, occasionally resulting in pathological fracture. Clinical manifestations of osteoporosis are the late consequences of abnormal bone formation and excessive regional overmodeling, compatible with the observation of increased osteoclast recruitment to bone surfaces in some areas.²¹ Cyclic infusion of bisphosphonate may offer partial relief of pain as it is a potent analgesic in painful bone diseases and improves bone mineral density. The use of bisphosphonates is limited by the increasing risks for rare complications such as bisphosphonate-related atypical fractures. The gradual disappearance of the diaphyseal widening in the metacarpals in long-surviving patient 5 exemplifies the effects of late progressive overmodeling of qualitatively abnormal cortical and trabecular bone. The qualitative degrading and dysfunction of soft connective tissue adversely affects not only periarticular and peri-osseous structures, but also bronchopulmonary elasticity and cardiac valve anatomy and function, ultimately contributing to lethal outcome.² Elucidation of the mechanisms by which this mutation leads to partial ‘uncoupling’ of the severe physical impact and the rather moderate cognitive disability in this specific intermediate ML will expand understanding of pathogenesis of ML beyond inadequately phosphorylated glycoproteins.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

ACKNOWLEDGEMENTS

Funding has been provided by Award Number U54NS065768 from the National Institute of Neurological Disorders And Stroke and the Office of Rare Diseases Research (ORDR) through the Lysosomal Disease Network Consortium, a part of the NIH Rare Diseases Clinical Research Network (RDCRN), and by the National Institute of General Medical Sciences (GM086524). Additional support has been provided by ISMRD, the International Advocate for Glycoprotein Storage Diseases, the National MPS Society, and by the Genetics Endowment of South Carolina.

- Cathey SS, Leroy JG, Wood T *et al*: Phenotype and genotype in mucopolipidosis II and III alpha/beta: a study of 61 probands. *J Med Genet* 2010; 478–488.
- Kobayashi H, Takahashi-Fujigasaki J, Takahiro F *et al*: Pathology of the first autopsy case diagnosed as mucopolipidosis type III $\alpha\beta$ suggesting autophagic dysfunction. *Mol Genet Metab* 2011; 102: 170–175.
- Kozłowski K, Lipson A, Carey W: Mild I-cell disease, or severe pseudo-Hurler polydystrophy in three siblings; further evidence for intermediate forms of mucopolipidosis II and III. Radiological features. *Radiol Med* 1991; 82: 847–851.
- David-Vizcarra G, Briody J, Ault J *et al*: The natural history and osteodystrophy of mucopolipidosis II and III. *J Paediatr Child Health* 2010; 46: 316–322.
- Thomas GH, Taylor HA, Reynolds LW *et al*: Mucopolipidosis 3 (Pseudo-Hurler polydystrophy): multiple lysosomal enzyme abnormalities in serum and cultured fibroblast cells. *Pediatr Res* 1973; 7: 751–756.
- Braulke T, Geuze HJ, Slot JW *et al*: On the effects of weak bases and momensin on sorting and processing of lysosomal enzymes in human cells. *Eur J Cell Biol* 1987; 43: 316–321.
- Stee R, Hullin R, Kudo M *et al*: A splicing mutation in the $\alpha\beta$ GlcNAc-1-phosphotransferase gene results in an adult onset form of mucopolipidosis III associated with sensory neuropathy and cardiomyopathy. *Am J Med Genet A* 2005; 132: 369–375.
- Reitman ML, Kornfeld S: UDP-N-acetylglucosamine: glycoprotein N-acetylglucosamine-1-phosphotransferase proposed enzyme for the phosphorylation of the high mannose oligosaccharide units of lysosomal enzymes. *J Biol Chem* 1981; 256: 4275–4281.
- Reitman ML, Kornfeld S: Lysosomal enzyme targeting. N-acetylglucosaminylphosphotransferase selectively phosphorylates native lysosomal enzymes. *J Biol Chem* 1981; 256: 11977–11980.
- Spranger JW, Brill PW, Poznanski AK: Bone Dysplasias. *An Atlas of Genetic Disorders of Skeletal Development*, 2nd (edn). New York: Oxford University Press 2002; pp 261–262.
- Pazzaglia UE, Beluffi G, Bianchi E *et al*: Study of bone pathology in early mucopolipidosis II (I-Cell disease). *Eur J Pediatr* 1989; 148: 553–557.
- Leroy JG, Cathey S, Friez MJ: *Mucopolipidosis II in: GeneReviews at GeneTests: medical genetics information resource (database online)*. Copyright. Seattle: University of Washington, 2012, pp 1997-2010. Available at: <http://www.genetests.org>
- Leroy JG, Cathey S, Friez MJ: *Mucopolipidosis III alpha/beta in: GeneReviews at GeneTests: medical genetics information resource (database online)*. Copyright. Seattle: University of Washington, 2012, pp 1997-2010. Available at: <http://www.genetests.org>
- Reitman ML, Varki A, Kornfeld S: Fibroblasts from patients with I-cell disease and pseudo-Hurler polydystrophy are deficient in uridine 5'-diphosphate-N-acetylglucosamine: glycoprotein N-acetylglucosaminyl-phosphotransferase activity. *J Clin Invest* 1981; 67: 1574–1579.
- Franke M, Braulke T, Storch S: Transport of the GlcNAc-1-phosphotransferase $\alpha\beta$ -subunit precursor protein to the Golgi apparatus requires a combinatorial sorting motif. *J Biol Chem* 2012; 288: 1238–1249.
- Varki AP, Reitman ML, Kornfeld S: Identification of a variant of mucopolipidosis III (pseudo-Hurler polydystrophy): a catalytically active N-acetylglucosaminylphosphotransferase that fails to phosphorylate lysosomal enzymes. *Proc Natl Acad Sci USA* 1981; 78: 7773–7777.
- Kudo M, Brem MS, Canfield WM: Mucopolipidosis II (I-cell disease) and Mucopolipidosis IIIA (classical pseudo-Hurler polydystrophy) are caused by mutations in the GlcNAc-phosphotransferase $\alpha\beta$ subunits precursor gene. *Am J Hum Genet* 2006; 78: 451–463.
- Amritraj A, Peake K, Kodam A *et al*: Increased activity and altered subcellular distribution of lysosomal enzymes determine neuronal vulnerability in Niemann-Pick type C1-deficient mice. *Am J Pathol* 2009; 175: 2540–2556.
- Nakanishi H: Neuronal and microglial cathepsins in aging and age-related diseases. *Ageing Res Rev* 2003; 2: 367–381.
- Tynnälä J, Sohar I, Sleat DE *et al*: A mutation in the ovine cathepsin D gene causes a congenital lysosomal storage disease with profound neurodegeneration. *EMBO J* 2000; 19: 2786–2792.
- Robinson C, Baker N, Hoffman P *et al*: The osteodystrophy of mucopolipidosis type III and the effects of intravenous pamidronate treatment. *J Inher Metab Dis* 2002; 25: 681–693.