Osteogenesis Imperfecta: Clinical Diagnosis, Nomenclature and Severity Assessment

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Manuscript Received: 31 October 2013; Manuscript Accepted: 12 February 2014

Recently, the genetic heterogeneity in osteogenesis imperfecta (OI), proposed in 1979 by Sillence et al., has been confirmed with molecular genetic studies. At present, 17 genetic causes of OI and closely related disorders have been identified and it is expected that more will follow. Unlike most reviews that have been published in the last decade on the genetic causes and biochemical processes leading to OI, this review focuses on the clinical classification of OI and elaborates on the newly proposed OI classification from 2010, which returned to a descriptive and numerical grouping of five OI syndromic groups. The new OI nomenclature and the pre-and postnatal severity assessment introduced in this review, emphasize the importance of phenotyping in order to diagnose, classify, and assess severity of OI. This will provide patients and their families with insight into the probable course of the disorder and it will allow physicians to evaluate the effect of therapy. A careful clinical description in combination with knowledge of the specific molecular genetic cause is the starting point for development and assessment of therapy in patients with heritable disorders including OI.

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Key words: osteogenesis imperfecta; fractures; collagen type I; heterogeneity; classification

INTRODUCTION

Osteogenesis imperfecta (OI) is the collective term for a heterogeneous group of connective tissue syndromes characterized primarily by liability to fractures throughout life. Since the first scientific description of OI in 1788 [Peltier, 1981; Baljet, 2002] the nomenclature and classification of OI has evolved substantially.

CLASSIFICATION AND NOMENCLATURE Osteogenesis Imperfecta 1979: The Original Sillence Classification

The present nosology and classification is based on the publication in 1979 by Sillence et al. [1979] entitled "*Genetic Heterogeneity in*

How to Cite this Article:

Van Dijk FS, Sillence DO. 2014. Osteogenesis imperfecta: Clinical diagnosis, nomenclature and severity assessment.

Am J Med Genet Part A 164A:1470–1481.

Osteogenesis Imperfecta". In this epidemiological and genetic study, 180 patients with OI were studied. OI patients were classified in four syndromes by primary clinical characteristics and pattern of inheritance namely (i) Dominantly inherited OI with blue sclerae, (ii) Lethal perinatal OI with radiographically crumpled femora and beaded ribs, (iii) Progressively deforming OI, and (iv) Dominantly inherited OI with normal sclerae. It is of note, that in the manuscript draft no numbers were given to the syndromes, which had been identified. The numbers OI types I–IV¹ were inserted in a table following a meeting with Dr. Victor McKusick who wanted to be able to put these syndromes into the computerized database, Mendelian Inheritance in Man (MIM). As such, the initial types I–IV reflected the order of appearance of the OI groups in the manuscript.

The four OI groups each displayed different modes of inheritance with autosomal dominant the predominant mode of inheritance for group I and IV and at least some families showed autosomal recessive inheritance for OI type II and III, indicating

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DOI 10.1002/ajmg.a.36545

¹In the "Nosology and classification of genetic skeletal disorders" Warrman et al. [2011] recommend using Arabic integers. We agree. However, since this introduction is addressing the historical aspects we have retained the Roman numerals where appropriate for clarity.

genetic heterogeneity in OI. OI type II was subsequently subdivided in OI type II-A, B, and C based on radiological features [Sillence et al., 1984]. In 1983, the first genetic cause of OI, an internal deletion in a collagen gene (COL1A1), was described in a patient with OI type II [Chu et al., 1983]. In the following years, mutations in the COL1A1 and COL1A2 genes encoding respectively the alpha1 and alpha2 chains of collagen type I were detected in all OI types. As such, the assumption of genetic (locus) heterogeneity in OI was largely abandoned in favor of allelic heterogeneity. It appeared that the specific type and position of the mutation (genotype) did affect the phenotype as had been proposed in the past [Caniggia et al., 1958; Smars, 1961]. However, some OI families remained without an identified genetic defect in the COL1A1/2 genes [Wallis et al., 1993]. In 2004, an expanded Sillence classification was published by Glorieux and Rauch [Rauch and Glorieux, 2004] adding OI types V-VII with an unknown genetic defect and presumed autosomal dominant inheritance (OI type V) and autosomal recessive inheritance (OI type VI and VII).

In 2006 the first autosomal recessive cause of OI type II was described i.e. CRTAP mutations [Barnes et al., 2006]. At present, a total of 17 genetic causes of OI have been described (see Table I, Fig. 1) with COL1A1/2 mutations still accounting for a large majority of OI patients, approximating 90% in populations of European origin [van Dijk et al., 2012]. Nomenclature revisions have accordingly seen the numbers of OI types being increased up to OI type XIV [Forlino et al., 2011] with the discovery of each new genetic defect. This is confusing in clinical practice since the newly added OI types, result in an OI classification in which the types are not mutually exclusive. OI types I-IV were defined because of specific clinical/radiological characteristics whereas the newer OI types (except for type V) were defined because they involved different gene loci with the clinical/radiological characteristics still being comparable within OI types II-IV [van Dijk et al., 2010].

Osteogenesis Imperfecta 2010: A New Ol Nomenclature

At the 2009 meeting of the International Nomenclature group for Constitutional Disorders ICHG of the Skeleton (INCDS) (Published as 2010 Nosology), a decision was finally made to group the known OI syndromes into five groups, that is, preserving the primary four groups and adding OI type V. The importance of the different genetic causes of the OI types was acknowledged by encapsulating the causative genes as subtypes of OI types I–V (Table I) [Warman et al., 2011].

The new nomenclature has also attempted to return to a descriptive grouping of syndromes (Table I) as was the case in the original description of the four OI types, which were defined because of specific clinical characteristics and inheritance pattern. While the ordering is unconventional to those who have used the numerical shorthand for the past 30 years, the order more closely reflects the historical sequence of discovery and has some phenotypic grading reflecting severity. It may be a surprise to some that the former syndrome of OI type IV is designated as Common Variable OI with normal sclerae. This phenotypic

variability within families was first noted by Ekman 1788 [Peltier, 1981] and the variability reinforced in the classics of OI in papers by Holcomb [1931] and the doctoral thesis of Seedorf [1949] which reported in detail the variability in families with different syndromes of OI.

The Nosology 2010 grouped OI with the Decreased Bone Density Group of Skeletal Dysplasias [Warman et al., 2011]. In addition to the syndromes with "brittle bones" and/or osteoporosis, encompassed by the descriptive groupings (previously numbered) of osteogenesis imperfecta syndromes, there is the large group of syndromes with decreased bone density, which have significant clinical overlap with the OI syndromes. These syndromes which are characterized by bone fragility and/or osteoporosis alone or with additional features, such as multiple contractures of the large joints as in Bruck syndromes 1 and 2 (Table II) have been included with the OI syndromes since 1992 and will also be included in the 2014 Revised Nosology. Premature aging syndromes in which fractures may be the first manifestation and precede noticeable hair loss, acro-osteolysis and skin aging have been in the main classified with the Acro-osteolyses Group of Disorders.

Because the recommended nosology is phenotypic, yet a numerical classification has been in use for over 30 years we have retained the concept of a short hand but adopted the use of Arabic numerals at this point to replace Roman numerals which were meant to imply distinct gene loci. Furthermore it was apparent from the original paper setting out a numerical nomenclature [Sillence et al., 1979] that the authors both in their discussion and tabulation had concluded that each of these phenotypic groups was likely to be genotypically heterogeneous.

SEVERITY GRADING IN OSTEOGENESIS IMPERFECTA SYNDROMES

In the years following the discovery of COL1A1/2 mutations in all OI types, the four OI types were often used in clinical practice to reflect severity with mild (OI type 1), lethal (OI type 2), severely deforming (OI type 3), and moderately deforming (OI type 4). Although the INCDS agreed to retain the Sillence classification as "the prototypic and universally accepted way to classify the degree of severity in Ol" [Warman et al., 2011], the need for internationally agreed criteria for grading severity between affected individuals was proposed and adopted, reflecting also the improved treatment possibilities (surgical, pharmacological and conservative) for patients with OI. The severity grading scale proposed here relies on clinical, historical data, fracture frequency, bone densitometry, and level of mobility (Table III). This severity grading was adopted for the POISE (Pediatric Osteogenesis Imperfecta Safety and Efficacy study) of Risedronate in osteogenesis imperfecta in 231 children ascertained from 22 investigators drawn from 11 countries [Munns and Sillence, 2013; Bishop et al., 2013]. The grading for the POISE study is modified here by the authors with addition of a general guideline to prenatal clinical and ultrasonographic findings. The scale will require further validation by collaboration between Centres of Expertise with sufficient patients and access to facilities for comprehensive assessment in order to further confirm and clarify its clinical utility.

TABLE I.	A Ne	ew Ol	Nomenclature	Combined	With	Causative	Genes	(A)	Phenotype	s With	Mild t	o Moderate	Severity,	(B)	Progressively
Deforming and Perinatally Lethal Phenotypes															

OI syndrome names	Туре	Gene	МІМ	Locus	Protein product	Inheritance
Non-deforming OI with blue sclerae	1	1. <i>COL1A1</i>	#166200	17q21.33	Collagen alpha-1(I) chain	AD
		2. <i>COL1A2</i>	#166200	7q22.3	Collagen alpha-2(I) chain	AD
Common variable Ol with normal sclerae	4	1. <i>COL1A1</i>	#166220	17q21.33	Collagen alpha-1(I) chain	AD
		2. <i>COL1A2</i>	#166220	7q22.3	Collagen alpha-2(I) chain	AD
		3. <i>WNT1</i> ^a	#615220	12q13.12	Wingless-type MMTV integration site family, member 1	AD
					Cartilage-associated protein (CRTAP)	
		1. CRTAP	#610682	3p22.3	Cyclophilin B (CyPB)	AR
		2. <i>PPIB</i>	#259440	15q22.31	Osterix	AR
		3. <i>SP7</i>	#613849	12q13.13	Plastin 3	AR
	-	1. <i>PLS3</i>		Xq23		XL
UI with calcification in interosseous membranes (B)	5	1. <i>IFIIM5</i>	#610967	11p15.5	Interferon-induced transmembrane protein 5	AD
Progressively deforming	3	1. <i>COL1A1</i>	#259420	17q21.33	Collagen alpha-1(I) chain	AD
5 5 5		2. <i>COL1A2</i>	#259420	, 7q22.3	Collagen alpha-2(I) chain	AD
		1. <i>BMP1</i>	#614856	8p21.3	Bonemorphogeneticprotein 1	AR
		2. CRTAP	#610682	3p22.3	Cartilage-associatedprotein (CRTAP)	AR
		3. <i>FKBP10</i>	#610968	17q21.2	Peptidyl-prolyl cis-transisomerase FKBP10	AR
		4. LEPRE1	#610915	1p34.2	Prolyl 3-hydroxylase 1 (P3H1)	AR
		5. <i>PLOD2</i>	#609220	3q24	Procollagen-lysine, 2-oxoglutarate	AR
		6. <i>PPIB</i>	#259440	15q22.31	5-dioxygenase 2	AR
		7. SERPINF1	#613982	17p13.3	Cyclophilin B (CyPB)	AR
		8. SERPINH1	#613848	11q13.5	Pigment-epithelium-derived factor (PEDF)	AR
		9. TMEM38B	#615066	9q31.1	Heat shock protein 47 (HSP47)	AR
		10.WNT1	#615220	12q13.12	Trimeric intracellular cation channel B (TRIC-B)	AR
		11.CREB3L1		11q11	Wingless-type MMTV integration site family, member 1	AR
					Old Astrocyte	AR
					Specifically induced substance (OASIS)	
Perinatally lethal Ol	2 ^b	1. <i>COL1A1</i>	#166220	17q21.33	Collagen alpha-1(l) chain	AD
		2. COL1A2	#166220	7q22.3	Collagen alpha-2(I) chain	AD
		1. CRTAP	#610682	3p22.3	Cartilage-associated protein (CRTAP)	AR
		2. LEPRE1	#610915	1p34.2	Prolyl 3-hydroxylase 1 (P3H1)	AR
		3. <i>PPIB</i>	#259440	15q22.31	Cyclophilin B (CyPB)	AR

^aSo far, 12 families with AR OI due to *WNT1* mutations have been described. Developmental delay was reported in affected individuals from three families. It is uncertain whether this is part of the clinical phenotype resulting from *WNT1* mutations [Fahiminiya et al., 2013; Keupp et al., 2013; Pyott et al., 2013]. A dominant *WNT1* mutation appeared to cause early onset osteoporosis [Keupp et al., 2013; Laine et al., 2013].

^bIn clinical practice subdivisions OI type II-A and OI type II-B are still in use. OI type II-A appears to be exclusively caused by heterozygous mutations in the *COL1A1/2* genes [van Dijk et al., 2010]. ^cIt has been reported that mutations in *PLOD2* may also result in progressively deforming OI [Puig-Hervás et al., 2012].

CLINICAL PRESENTATIONS AND FEATURES OF OSTEOGENESIS IMPERFECTA Clinical Presentations and Features of OI in General

Primary feature: liability to fractures and osteoporosis. While liability to fractures throughout life is the single most important clinical feature, experience with families with OI type 1 indicate that perhaps 10% of affected individuals have not had a long bone fracture during childhood [Sillence, 1980]. However, newer

techniques for measuring bone density, such as dual energy X-ray absorptiometry (DXA) of the skeleton [Lu et al., 1994] and/or more recently peripheral quantative computerized tomography (pQCT) [Gatti et al., 2003; Folkestad et al., 2012] of forearm and leg, frequently reveal significantly reduced bone density in a least one area of the skeleton in those individuals who by formal genetic analysis have OI (Table III).

Net bone fragility is the final result of contributions from primary bone fragility and the secondary fragility resulting from osteoporosis. Osteoporosis develops in the majority of patients with OI. The finding of elevated serum and urine markers of bone



FIG. 1. Overview of collagen type I biosynthesis. Collagen type I consists of two α 1-chains and one α 2-chain. After translation, pro- α 1-chains and pro- α 2 chains are processed in the rough Endoplasmic reticulum (rER). These chains have to align in order to start the folding process of (pro)collagen type I into a triple helix. The next step is alignment of the three chains in order to commence folding into a triple helical structure. During this folding process, post-translational modification by specific proteins takes place. The genes encoding proteins involved in post-translational modification and in which mutations have been reported to cause 0I, are depicted in this figure. After transport of procollagen type I to the Golgi complex and following exocytosis into the extracellular matrix, cleavage of the C-and N-propeptides results in formation of collagen type I. Subsequently, cross-linking of collagen type I molecules leads to formation of fibrils. Multiple collagen type I fibrils form into collagen fibers, important constituents of bone.

turnover in patients with OI considered along with the findings of bone histomorphometry, is best explained by a combination of increased bone formation and increased bone resorption [Rauch and Glorieux, 2004]. The net effect is a small progressive bone loss since bone resorption is often greater than bone formation, with immobilization also exerting a negative effect on bone formation. Bisphosphonate treatment aimed at reduction of osteoclast activity, is initiated in many children with OI after careful assessment by the treating physician. In that regard cyclical treatment with intravenous bisphosphonates has become the gold standard for treatment of children with moderate to severe OI. A very recent randomized, double-blind, placebo-controlled trial of oral Risedronate in children with OI, including a large proportion of

Disorder	МІМ	Gene	Locus	Protein Product	Inheritance
Familial doughnut lesions of skull	#126550	Unknown	Unknown	Unknown	AD
Geroderma osteodysplasticum	#231070	GORAB	1q24.2	RAB6-interacting golgin	AR
Gnathodiaphyseal dysplasia (osteopenia with radiolucent lesions of the mandible	#166260	ANO5	11p14.3	Anoctamin-5	AD
Hajdu–Cheney syndrome	#102500	NOTCH2	1p12-p11	Neurogenic locus notch homolog protein 2	AD
ldiopathic juvenile osteoporosis ^a	259750	Unknown	Unknown	Unknown	Unknown
Infantile hypophosphatasia	#241500	ALPL	1p36.12	Alkaline phosphatase, tissue- nonspecific isozyme	AR
Ol with congenital joint contractures type 1 (Bruck syndrome type 1) ^b	#259450	FKBP10	17q21.2	Peptidyl-prolyl cis-trans isomerase FKBP10	AR
Ol with Congenital Joint Contractures Tupe 2 (Bruck syndrome type 2)	#609220	PLOD2	3q24	Procollagen-lysine, 2-oxoglutarate 5-dioxygenase 2	AR
Osteogenesis imperfecta with craniosynostosis, ocular proptosis, hydrocephalus, and distinctive facial features (Cole-carpenter)	112240	Unknown	Unknown	Unknown	AD
Osteoporosis pseudoglioma	#259770	LRP5	11q13.4	Low density lipoprotein-related protein 5	AR
Primary osteoporosis	#259770	LRP5	11q13.4	Low density lipoprotein-related protein 5	AD
Spondylocular syndrome (osteoporosis, cataracts and retinal dysplasia)	605822	Unknown	Unknown	Unknown	AR

TABLE II. Syndromes With Pre-and/or Postnatal Phenotypic Features Overlapping OI

^aGenetically heterogeneous. A small percentage of patients with idiopathic juvenile osteoporosis have been classified as primary genetic osteoporosis with heterozygous mutations in the low density lipoprotein related protein 5 (*LRP5*). As such, these latter usually have a milder bone disorder than patients with osteoporosis pseudoglioma who have homozygous or compound heterozygous mutations in *LRP5* and severe osteoporosis with visual disability [Hartikka et al., 2005].

^bRecently, recessive mutations in FKBP10 have been described to cause (a) recessive OI most closely resembling OI type III or (b) Bruck syndrome type I [Schwarze et al., 2013].

more mild to moderately affected children, demonstrated a significant reduction in fracture risk, thus extending the therapeutic benefits of this therapy in children with OI [Bishop et al., 2013].

Associated features in general. Associated features in some affected individuals, but not others, include distinct blueness of the sclerae, young adult onset hearing loss, dentinogenesis imperfect (DI), joint hypermobility, short stature, and progressive skeletal deformity. Cardiovascular complications such as valvular dysfunction and aortic root dilatation have been reported in adult OI patients, more often in patients with OI type 3 [Radunovic et al., 2011]. Several associated features are more elaborately described below.

Blue Sclerae. Several major reviews and at least one monograph [Smars, 1961; Sillence et al., 1979; Sillence et al., 1993] concluded that in patients with "blue sclerae" the color of the sclerae is similar to Wedgewood blue in hue and is so very distinctive that the sclerae appear painted. When "blue sclerotics" are present, they remain distinctly blue throughout life.

Berfenstam and Smårs in a population based study [1956] showed that there were statistically significant differences in patterns of phenotypic symptoms and musculoskeletal complications between two groups of patients with OI, those with blue-grey sclerae and those with normal sclerae. Data from a cohort of 95 patients with OI type 1 and OI type 4 in the 1979 Victorian population study were reanalysed to confirm that finding [Sillence et al., 1993]. Attention was also drawn to the misconception that the bluegray sclerae in OI type 1 are due to the thinning of the sclerae. Eichholtz and Muller [1972] had reported that overall thickness of the sclerae in OI type 1 was normal and there was increased electron dense granular material between scleral collagen fibers. It was proposed that in the pathogenesis of OI type 1, the hearing impairment, easy bruising and possibly the marked joint hypermobility would be best explained by secondary dysregulation of connective tissue composition. There is further evidence that the high prevalence of premature termination/nonsense/splicing mutations which cause the OI type 1 phenotype are associated with alterations in matrix composition [Byers and Cole, 2002].

Dentinogenesis imperfecta. Dentinogenesis imperfecta produces a distinctive yellowing and apparent transparency of the teeth, which are often worn prematurely or broken. Some teeth may have a particularly greyish hue. Radiologic studies of affected teeth show that they have short roots with constricted corono-radicular junctions [Bailleul-Forestier et al., 2008].

Secondary deformations. Skeletal deformities such as scoliosis and basilar impression are regarded as secondary deformations rather than primary malformations. Although the absence of deformity of long bones has been advanced as a diagnostic criterion, the presence of deformity seems at least partly significantly influenced by quality of care. In developing countries, deformity may be evidence of sub-optimal care reflecting lack of primary care services



for managing fractures, rather than evidence of an intrinsic process of bone deformation.

Non-Deforming OI With Blue Sclerae—OI Type 1

OI type 1 is characterized by increased bone fragility, which is usually associated with low bone mass, distinctly blue-gray sclerae, and susceptibility to conductive hearing loss commencing in adolescence and young adult life. Deformity of long bones or spine is uncommon and where scoliosis develops it is commonly an idiopathic scoliosis. OI type 1 is the most common variety of OI in European derived communities and has a birth prevalence in the order of 1:25,000 live births and a similar population frequency [Steiner et al., 2013]. Fracture frequency and usually mild long bone and spine deformity mean that it is generally perceived to be of mild severity but occasionally it is moderately severe, particularly when DI is present [Paterson et al., 1983].

DI is observed in some families with this trait and not others. Paterson and colleagues showed that patients with OI type 1 and DI are more likely to have fractures at birth (25% vs. 6%) than those without DI. Furthermore, patients with OI type 1 and DI have a higher fracture frequency, more severe short stature, and more skeletal deformity. Both subgroups have a similar frequency of joint hypermobility, bruising, deafness, and joint dislocations [Paterson et al., 1983].

Hearing impairment resulting from both conductive and sensorineural loss is detectable in over 50% of patients with OI type 1 by 40 years of age [Kuurila et al., 2002; Swinnen et al., 2011]. Vertigo is a troublesome symptom in many people with OI including OI type 1 [Kuurila et al., 2003].

Families with autosomal dominant inheritance and variable expressivity have been reported in many studies. Penetrance for blue sclerae is close to 100% but frequency of clinical fractures is only 90–95% [Smars, 1961; Sillence et al., 1979].

Common Variable OI—OI Type 4

These patients have recurrent fractures, osteoporosis and variable degrees of deformity of long bones and spine but normal sclerae. The sclerae may be bluish at birth but the blue tinge fades during childhood. Hearing impairment is not often encountered. Posterior fossa compression syndromes due to basilar impression with elevation of the floor of the posterior cranial fossa are increased in prevalence. Patients with OI type 4 who have DI have a five times higher relative risk for basilar impression [Sillence, 1994]. Some 30% of patients with OI type 4 have basilar impression on screening but only 16% of these are symptomatic [Sillence, 1994]. Common variable osteogenesis imperfecta with normal sclerae shows occasionally autosomal recessive [van Dijk et al., 2010] and X-linked inheritance [van Dijk et al., 2013] but it is usually inherited as an autosomal dominant disorder (Table I). Severity is highly variable within families. It is not uncommon to find families where there are many affected with mild OI but a few individuals in the same family with moderately severe OI [Holcomb, 1931; Seedorf, 1949].

Progressively Deforming OI–OI Type 3

Individuals with OI type 3 usually have newborn or infant presentation with bone fragility and multiple fractures leading to

progressive deformity of the skeleton. They are generally born at or near term and have normal birth weight and often normal birth length, although this may be reduced because of deformities of the lower limbs at birth. Although the sclerae may be blue at birth, observation of many patients with this syndrome documents that the sclerae become progressively less blue with age [Sillence et al., 1986]. Persisting blue sclerae are usually an indication of nonsense or frameshift mutations in type 1 collagen genes characteristic of non-deforming OI type 1 whereas patients with the various autosomal recessive disorders will usually have grey-white sclerae [Byers and Pyott, 2012]. All patients have poor longitudinal growth and fall well below the third centile in height for age and sex. Progressive kyphoscoliosis develops during childhood and progresses into adolescence. Hearing impairment has not been reported in children with this syndrome but hearing loss is more frequent in adults. DI is a variable feature.

At birth, radiographic studies show generalized osteopenia and multiple fractures. Bowing and angulation deformities exist to a variable degree with frequent over-modeling of the shafts of the long bones. Within weeks to months, in some infants, undermodeling of the shafts of long bones results in a "broad-bone" appearance. From several years of age, metaphyses develop increasing density and irregularity. These metaphyseal changes designated a "pop-corn" appearance may evolve only to resolve completely after puberty. The ribs are thin, osteopenic, and progressively crowded as platyspondyly increases. The skull shows multiple Wormian bones, although these may not be evident until several weeks to months of age [Sillence et al., 1979; Sillence et al., 1986; Spranger et al., 2003; van Dijk et al., 2011].

In the past, approximately two-thirds of the patients died by the end of the second decade. Death usually resulted from the complications of skeletal chest wall deformity including kyphoscoliosis, pulmonary hypertension, and cardio-respiratory failure. With the present therapeutic options, specifically bisphosphonate treatment with cyclic intravenous Pamidronate [Glorieux et al., 1998] commenced in infancy, it can be expected that today the majority of patients with OI type 3 will survive into adult life. Several studies demonstrate that centers of expertise that manage children with severe OI, achieve very reduced fracture frequency and near normal growth velocity in infants commenced on cyclic intravenous pamidronate by 3 years of age [Plotkin et al., 2000; DiMeglio et al., 2004; Munns et al., 2005; Astrom et al., 2007]. A recent publication confirmed that treatment appears to be well tolerated and associated with an increase in bone density, reduced fracture frequency and improved vertebral shape [Alcausin et al., 2013].

Perinatally Lethal OI Syndromes—OI Type 2

The skeletal, joint, and extraskeletal features of this group of fetuses and children are extremely severe. Perinatal lethality is an outcome rather than a diagnostic feature. Fetuses detected at 18–20 weeks gestation have short crumpled long bones, bowing or angulation deformities of long bones and marked deficiency of ossification of facial and skull bones. At this early gestation, there may be few rib fractures but with each month in utero there is progressive fracturing of ribs resulting in the continuously beaded appearance combined with crumpled (accordion-like) long bones that is characteristic of the extremely severe end of the spectrum represented by OI type 2 (OI type 2-A) [Sillence et al., 1984]. In our experience, treatment with cyclic intravenous pamidronate is not indicated as bone formation is so impaired and joint restriction so severe there is virtually no chance of any normal childhood life experience. Pain relief with simple analgesics or subcutaneous morphine is particularly valuable, improving comfort and breathing. Among the extraskeletal features in OI type 2, neuropathological findings such as brain migrational defects and/or white matter changes have been reported in a limited number of cases [Emery et al., 1999]. Some babies have a phenotype which is a little less severe with fewer rib fractures (OI type 2-B) [Sillence et al., 1984] and as such they can show overlap with OI type 3 [Spranger, 1984]. Rarely these babies survive, even to adult life and can be "rescued" with treatment with cyclic intravenous pamidronate.

In developed countries, many or most children with OI type 2 are at present diagnosed prenatally (by ultrasound and DNA analysis), often resulting in termination of pregnancy. Mean birth length and weight are less than the fiftieth centile [Sillence et al., 1984]. The thighs are held abducted and in external rotation. The chest is small for gestation and respiratory excursion may be depressed because of the pain from multiple rib fractures and the abnormal biomechanical properties of semicontinuous beading from fracture callus along each rib in the most severely affected. Several clinical features suggest that newborns with OI type 2 are in constant pain. They may have excessive perspiration, pallor, show anxiety at being touched and move their limbs very little because of multiple fractures. One-fifth are stillborn and 90% die by 4 weeks of age [Sillence et al., 1984].

OI With Calcification in Interosseous Membranes—OI Type 5

OI type 5 with moderate to severe bone fragility was originally defined by Battle and Shattock [1908] as a type of OI with progressive calcification of the inter-osseous membranes in the forearms and legs. Independently it was identified by increased propensity to develop hyperplastic callus. The syndrome was delineated in some detail by Bauze et al. [1975], who observed that 10% of patients with moderate to severe OI and normal sclerae, had OI type 5 [Bauze et al., 1975]. In a histomorphometric study of moderately severe OI type 4, 7 of 26 cases (25%) were detected with abnormal bone histomorphometry which is characteristic of OI type 5 [Glorieux et al., 2000]. In clinical studies it accounts for approximately 5% of individuals with OI seen in a hospital setting.

Calcification of the inter-osseous membrane in the forearms is observed from early in life, which leads to restriction of pronation and supination, and eventual dislocation of the radial heads. The sclerae are white and DI and Wormian bones are not present. Those affected tend to have higher serum alkaline phosphatase values and have an increased risk of developing hyperplastic callus following a fracture or orthopaedic surgery. A distinct pathogenesis is further supported by characteristic bone histomorphometry which shows coarse mesh-like lamellation which distinguishes OI type 5 from OI type 4 [Glorieux et al., 2000]. Hyperplastic callus is a rare medical emergency occurring in patients with OI type 5. This is characterized by massive callus with swelling and pain at the site of a fracture, which may be as minor as a stress fracture. Prompt use of indomethacin, an anti-inflammatory COX-1 and COX-2 prostaglandin inhibitor has been recommended to avert progress although of the callus although a randomized clinical trial has not been reported [Glorieux et al., 2000; Cho and Moffat, 2014].

MOLECULAR GENETICS OF OI

Currently, more than 1,000 heterozygous *COL1A1/2* mutations have been identified (https://oi.gene.le.ac.uk, accessed April 1 2013) [Dalgleish, 1997, 1998]. Mutation type and position influence the phenotype and as such genotype–phenotype relations exist to some extent.

Autosomal Dominant OI (OI Types 1-5)

In the majority of affected individuals from European descent, OI types 1–4 result from heterozygous mutations in the *COL1A1/2* genes encoding respectively the alpha1 and alpha2 chains of collagen type I (Fig. 1). The biosynthesis of collagen type I has been depicted in Table III. Sibling recurrence without an affected parent may occur due to gonadal mosaicism for heterozygous dominant mutations in one of the parents [Byers and Cole, 2002].

Patients with OI type 1 and sometimes OI type 4 have an approximately 50% reduction (quantitative or haploinsufficiency effect) in the synthesis of type 1 procollagen often due to heterozygous mutations in one COL1A1 allele (nonsense, frameshift, and splice site alterations) leading to mRNA instability and haploinsufficiency. Other causes are deletions of the whole COL1A1 allele or substitutions for glycine by small amino acids (cysteine, alanine, and serine) near the amino-terminal ends of the triple helical domains in either one COL1A1 or COL1A2 allele [van Dijk et al., 2012]. Notwithstanding the 50% reduction in collagen synthesis from individual cells, these patients have above average new bone formation, the result of homeostatic mechanisms, which increase the number of bone forming units. This increased new bone formation is linked to increased bone turnover so that the net effect is a small annual bone loss, which is exaggerated if there is immobilization because of fractures or pain [Rauch and Glorieux, 2004].

The majority of cases of OI type 2–4 in North America and Europe are dominantly inherited and most cases are due to heterozygous *COL1A1/2* mutations that result in substitutions for glycine. In general, glycine substitutions near the carboxyl-terminal end appear to result in the severest phenotype. Less common mutations include splice site alterations, insertion/deletion/duplication events that lead to in-frame sequence alterations and variants in the carboxyl-terminalpropeptide coding-domains [van Dijk et al., 2012] The heterozygous mutations disrupt triple helical assembly of type I collagen polypeptides, resulting in overprocessing by the enzymes responsible for post-translational modification of (pro) collagen type I and consequently production of abnormal collagen type I. This post-translational over-modification is demonstrable by SDS–polyacrylamide gel electrophoresis. The intertwining of mutated and normal collagen type I chains result in production of abnormal collagen type I protein, which is rapidly degraded (dominant-negative effect).

Recently, the genetic cause of OI type 5 has been elucidated in two independent publications [Cho et al., 2012; Semler et al., 2012] and consists of a heterozygous C>T transition in the 5'UTR (untranslated region) of *IFITM5* (c.-14C>T). *IFITM5* encodes Interferon induced transmembrane protein 5, the expression of this protein has been shown to peak during osteoblast formation in the early mineralization stage in mice and rats [Hanagata et al., 2011].

Autosomal Recessive OI (OI types 2-4)

A severe, autosomal recessive form of OI type 3 with a comparatively high frequency had already been recognized in the past in the black populations of southern Africa [Wallis et al., 1993] (Table III). Nowadays, it is also known that a founder mutation in LEPRE1 is carried by 1.5% of West Africans and 0.4% of African Americans [Cabral et al., 2012]. Recessive mutations in genes involved in collagen type I biosynthesis and post-translational modification have been identified in OI types 2-4 in the last 6 years. These were recently reviewed in depth [Byers and Pyott, 2012]. The recessive mutations concern genes encoding proteins involved in collagen type I biosynthesis, can be subdivided into (i) an enzymatic complex responsible for 3-prolyl hydroxylation of one specific residue (P986) in the alpha1 chain [van Dijk et al., 2012] and probably for initiating chain alignment and helical folding [Pyott et al., 2011] (CRTAP, LEPRE1, PPIB); (ii) quality control check of the collagen triple helix (SERPINH1, FKBP10); (iii) late processing of folded (pro) collagen type I chains i.e. hydroxylation of lysine residues in triple helical telopeptides important for collagen type I cross-linking in bone [van Dijk et al., 2012] (PLOD2, FKBP10) and cleavage of the C-propeptide (BMP1) [Martínez-Glez et al., 2012] (Fig. 1).

Furthermore, recessive mutations have been reported in *SP7* encoding Osterix, an osteoblast specific transcription factor, in *SERPINF1* possibly involved in bone formation and remodeling [van Dijk et al., 2012] and in *TMEM38B* encoding a trimeric intracellular cation channel [Shaheen et al., 2012; Volodarsky et al., 2013].

The recent delineation of mutations in WNT1, encoding a signaling peptide involved in osteoblast differentiation and proliferation [Fahiminiya et al., 2013; Keupp et al., 2013; Laine et al., 2013] and the interaction with FRIZZLED and its coligand LRP5, in which mutations in the latter are known to result in patients with severe syndromic OI, predict that further study of patients with severe OI from endogamous populations will uncover mutational mechanisms in the subsequent steps of the WNT-Beta Catenin signaling pathway. Most recently, a homozygous deletion of CREB3L1 was identified in a family with a severe progressively deforming OI phenotype. CREB3L1 encodes the ER-stress transducer OASIS that has been shown in a murine model to bind to the osteoblast-specific UPRE (unfolded protein response element) regulatory region in the Collal promotor. This finding expands the genetic heterogeneity in OI and illustrates the role of ER-stress in the pathophysiology of OI [Symoens et al., 2013].

Pathogenic mutations found in recessive genes (Dalgleish, R: Osteogenesis Imperfecta Variant Database (https://oi.gene.le.ac.uk,

accessed April 1 2013), are mostly homozygous or compound heterozygous loss-of-function mutations that result in two null alleles with severely decreased or no production of normal protein.

X-linked Ol

X-linked inheritance of osteoporosis and fractures had been reported only once in the thesis of D. Sillence (Pedigree 41, Appendix) [Sillence, 1980]. Recently, loss-of-function mutations in PLS3 encoding plastin-3 were discovered as a cause of one form of X-linked osteoporosis with fractures [van Dijk et al., 2013]. In hemizygous men, pathogenic mutations in PLS3 were associated with osteoporosis and osteoporotic fractures of the axial and appendicular skeleton usually developing in childhood. The clinical picture in heterozygous female members was variable and ranged from normal bone mineral density and an absence of fractures to early-onset osteoporosis. No extraskeletal features of OI were present in affected men, but the phenotype is indistinguishable in many patients with other types of OI, it would probably fit best in the common variable OI (OI type 4) group, of whom less than 50% of patients have features such as Wormian bones in the skull and the sclerae are normal in hue, bluish in childhood and fading to normal adult hue.

CONCLUSION

From a medical geneticist point of view, the core principle is phenotyping of individuals (dysmorphology) and the study of these families with regard to inheritance pattern and phenotypic variability. The OI classification from 1979 is a classic example of the importance and possibilities of dysmorphology since it led to the delineation of four OI syndromes based on clinical/radiological features and inheritance, in combination with the assumption that OI was genetically heterogeneous, which was confirmed many years later by molecular genetic studies.

At present time, it has been postulated that molecular techniques such as Next-Generation Sequencing will decrease the need for phenotyping. However, the new OI nomenclature and the Severity Grading Scale described in this paper, emphasize the importance of phenotyping in order to diagnose, classify and assess severity of OI. This will provide patients and their families with insight into the probable course of the disorder and it will allow physicians to evaluate the effect of therapy. A careful clinical description in combination with knowledge of the specific molecular genetic cause is the starting point for development and assessment of therapy in patients with heritable disorders including OI. The latter is the biggest challenge we face in the upcoming decade(s).

ACKNOWLEDGMENTS

Our teachers Professor David Danks, Murdoch Institute, Victoria (deceased), David L. Rimoin, Cedars Centre for Medical Genetics and International Skeletal Dysplasia Register (deceased), Ralph Lachman, MD, professor and former Chief of Paediatric Radiology Harbor-UCLA medical centre, are acknowledged for their guidance over several decades of research into the clinical genetic and molecular heterogeneity of osteogenesis imperfecta syndromes. When it comes to development of pharmacological therapy for OI, Professors DeVogaeler, Eva Astrom, and Francis Glorieux and colleagues have been of great importance in advancing a therapeutic praxis. In the fields of collagen biochemistry and molecular biology, Peter Byers, MD and his collaborator Dr. Ulrike Schwarze and their many colleagues at Centre for Collagen Research Seattle Washington, Hans-Peter Bachinger, PhD and colleagues, Shriners Portland Oregon, Joan Marini, MD, PhD and colleagues, Bethesda, Maryland, Dr. Raymond Dalgleish in Leicester, United Kingdom and Gerard Pals, PhD in Amsterdam, the Netherlands, have made outstanding contributions in taking the field of osteogenesis imperfecta forward. Of course many others, not specifically named must be acknowledged for their contribution to the field of OI.

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