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The limb-girdle muscular dystrophies- diagnostic strategies.

Kate Bushby¹, Fiona Norwood², Volker Straub¹.

1. Institute of Human Genetics,

International Centre for Life,

Central Parkway,

Newcastle upon Tyne

NE1 3BZ

Kate.bushby@ncl.ac.uk

0191 241 8757

0191 241 8799

(Professor Bushby is corresponding author)

2. King's Regional Neuroscience Centre

9th Floor Ruskin Wing

King's College Hospital

Denmark Hill

London

SE5 9RS

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Introduction

The limb-girdle muscular dystrophies are a group of disorders which were historically grouped together because of the shared clinical feature of predominant involvement of the "limb-girdle" (pelvic and shoulder) musculature. These clinical findings are not unique to the disorders within the "LGMD" classification- disorders such as Becker muscular dystrophy, late onset spinal muscular atrophy, myotonic dystrophy type 2, Bethlem myopathy and Pompe's disease may all show overlapping clinical features, and rarer disorders such as congenital myasthenic syndromes may also need to be considered in the differential diagnosis. Conversely, the LGMD classification also contains genetically defined disorders where there is a broad heterogeneity of presentation and muscle involvement. This is particularly true for some of the dominant disorders in the LGMD classification, such as LGMD1A where myotilin mutations are associated also with the disorders known as the myofibrillar myopathies, LGMD1B where lamin A/C mutations can also cause a wide range of different diseases, and LGMD1C where the clinical presentation includes hyperCKaemia, rippling muscle disease, myalgia and distal myopathy. The molecular basis of the diseases is also highly heterogeneous. The process of gene identification in LGMD has involved a combination of linkage and candidate gene analysis, resulting in a gene and protein based classification which includes three known genes causing an autosomal dominant LGMD [1] and eleven known genes causing autosomal recessive LGMD [2, 3] (see table 1). The genes and proteins causing the diseases in these groups show a huge range of localisation across the muscle fibre, from sarcolemma to nuclear envelope, with functions ranging from structural to enzymatic. As the current state of our understanding of the pathogenesis of many of these disease groups is dealt with elsewhere in this issue, this article will

concentrate on the diagnostic process by which these disorders can be defined. Such definition of a precise diagnosis is increasingly allowing directed management for these diseases by the ability to predict specific complications such as those of the cardiac or respiratory systems, and in the future will be the starting point for specific gene and protein based therapies.

History and examination

LGMD was first designated a disease group through the need to distinguish a set of patients with predominantly proximal muscle weakness and a dystrophic pattern on muscle biopsy from the more classically recognised "Duchenne/ Becker" and facioscapulohumeral muscular dystrophies, both of which groups are much more common than LGMD. The key hallmarks of disease remain the marked weakness of the pelvic and shoulder girdle muscles, but as the different disease entities within the group have become better defined based on their distinct genetic basis, accompanying clinical features may help to distinguish subtypes of LGMD. Particular features which need to be sought include muscle hypertrophy or atrophy, scapular winging, muscle rippling and contractures. While the history will in many cases be a relatively nonspecific story of progressive proximal weakness, with onset at any age, there are some clues in the history of onset which can be very suggestive of a particular diagnosis. This is particularly the case in dysferlinopathy where the history of relatively sudden onset in the late teens, often with early inability to stand on tiptoe, and sometimes with sudden onset of calf pain and swelling may be almost pathognomonic for the diagnosis [4]. This group of patients in particular may be misdiagnosed as myositis, with the inappropriate use of steroids, to which the patients do not appear to be responsive. In some of the autosomal dominant types of LGMD the family history

may be particularly informative. For example, the presence in the index case or wider family of respiratory failure, dysarthria or cardiomyopathy may help to suggest LGMD1A [5]. History of arrhythmia or contractures can be suggestive of laminopathy or LGMD1B [6], while a history of muscle rippling or myalgia may be suggestive of LGMD1C [7] (see table 2).

Associated complications

Muscular dystrophies are often multisystem disorders. Recognition of the particular type of muscular dystrophy can help to assign a more precise risk of complications. Complications which may be life-threatening and where appropriate management may have a positive impact on length and quality of life include especially those affecting the cardiovascular and respiratory systems. These are present at different frequencies within the various disorders in the LGMD group. Cardiovascular complications may include the development of a progressive cardiomyopathy or arrhythmias, while respiratory complications may or may not have significant diaphragmatic weakness as a clear component of the problem. These complications need to be sought specifically and appropriate management instituted for any problems detected [8, 9] (see table 2).

The diagnostic process (see figure 1)

None of the different types of LGMD can be distinguished on basic muscle histology or electrophysiology, though electromyography may be useful in distinguishing this myopathic group of diseases from myotonic dystrophy type 2 (PROMM, DM2) which may be clinically similar, and the muscle biopsy may be crucial in distinguishing LGMD from Pompe's disease or other metabolic syndromes. The serum creatine kinase is variably elevated in the different disorders and this may serve as a useful

indicator to subdivide the different types of LGMD, though it is no more than a guide. As a rule of thumb, the autosomal recessive types of LGMD are typically associated with a much higher level of creatine kinase elevation than the dominant forms of LGMD, with the exception of LGMD1C. Diagnosis therefore relies on a combination of clinical assessment, specialised muscle immunoanalysis and genetic testing. In informative families, linkage analysis may be useful to identify the gene involved. Otherwise, protein analysis can be used to pinpoint the likely gene primarily involved, though often complex patterns of secondary protein involvement in addition to the primary defect can complicate this process. A wide range of antibodies needs to be applied and frequently both immunolabelling of a muscle biopsy sample needs to be supplemented by immunoblotting. This is summarised in table 3. Combining information from the clinical and biopsy details therefore allows a diagnostic algorithm to be suggested and this is presented in figure 1. In part 1 of the algorithm the features which may suggest diagnosis of either LGMD2I or LGMD1B (neither of which can be definitively diagnosed by muscle biopsy) are described. Parts 2-4 describe the interpretation of the muscle biopsy findings together with clinical pointers which may lead to a suggestion of a genetic diagnosis which needs to be

Conclusions.

confirmed by testing.

Achieving a precise diagnosis in all patients will become increasingly important as new specific therapies are developed. The LGMDs continue to present a complex challenge for diagnosis, but the recognition of the different subgroups within this broad and heterogeneous classification already allows better strategies for management and genetic counselling. As diagnostic testing begins to be

systematically applied in different countries, the epidemiology of these diseases can increasingly be understood, and a variable pattern of disease frequency can be discerned. Therefore the future development of diagnostic testing and indeed therapeutic studies will need to take these regional differences into account.

Particularly striking is the predominance of LGMD2I due to the common C826A mutation in Northern Europe [10], compared to a relatively higher proportion of LGMD2A and LGMD2B cases in Southern Europe. A highly focussed approach to LGMD diagnostics is already improving management and is going to become increasingly important in the future.

Legends to tables and figures:

Table 1: The limb-girdle muscular dystrophies: classification of the entities for which the genetic basis is known.

Table 2. Key clinical features and complications of the limb girdle muscular dystrophies

Table 3. Diagnostic features: creatine kinase and muscle biopsy findings.

Figure 1:

A diagnostic algorithm for achieving diagnosis in LGMD based on clinical, creatine kinase, muscle biopsy and molecular genetic testing. CK= creatine kinase, CK levels are given as a general guide to levels of CK likely at diagnosis, but this can vary from patient to patient and with disease progression. CK level A= normal or mild elevation (<10x normal) CK level B= moderate elevation (8- 40x normal) CK level C extreme elevation (>40x normal). Other abbreviations: ADEDMD- autosomal dominant Emery Dreifuss muscular dystrophy, EDMDX- X linked EmeryDreifuss muscular dystrophy, PIRCs: percussion induced repetitive contractions.

Table 1: The limb-girdle muscular dystrophies: classification of the entities for which the genetic basis is known.

Type of	Gene/	Protein localisation/ function	Other phenotypes	References
LGMD	protein	(if known)		
	involved			
LGMD1A	Myotilin	Sarcomeric localisation	Myofibrillar myopathy	[11, 12]
LGMD1B	Lamin A/C	Nuclear envelope	Many- see chapter	[13, 14]
LGMD1C	Caveolin 3	Plasma membrane, caveolae	Dilated cardiomyopathy	[15, 16]

LGMD2A	Calpain 3	Cytoplasmic: proteolytic	[17, 18]	
		enzyme		
LGMD2B	Dysferlin	Predominantly plasma	Miyoshi Myopathy,	[19, 20]
		membrane: membrane repair	distal muscular	
			dystrophy with anterior	
			tibial involvement	
LGMD2C-F	Alpha, beta,	Part of dystrophin glycoprotein	Delta sarcoglycan:	[21, 22]
	gamma, delta	complex in plasma membrane	dilated cardiomyopathy	
	sarcoglycan	5		
LGMD2G	Telethonin	Sarcomeric localisation		[23]
LGMD2H	TRIM32	Ubiquitin ligase	Sarcotubular myopathy,	[24, 25]
			Bardet Biedl Syndrome	
LGMD2I	FKRP	Probable glycosyl transferase:	Congenital muscular	[26, 27]
		substrate alpha dystroglycan	dystrophy MDC1C and	
			intermediate phenotypes	
LGMD2J	Titin	Component of sarcomere	Heterozygotes have	[28]
			distal myopathy. Other	
		P	mutations associated	
	20		with cardiomyopathy	
LGMD2K	POMT1	Glycosyl transferase	Walker Warburg	[2]
	X		syndrome and	
			intermediate phenotypes	

Table 2. Key clinical features and complications of the limb girdle muscular dystrophies

Type of	Age at	Key features	Key features	Complications	Specific surveillance and treatment
LGMD	onset	from history	from examination		
LGMD1A	Usually	May be	May be distal	Cardiomyopathy,	Cardiac echo, ECG +/- Holter, FVC in sitting and lying.
	adult-	dysarthria, tight	muscle	arrhythmia, respiratory	Treatment with pacing, respiratory support
	hood	TAs, swallowing difficulties	involvement	failure	
LGMD1B	Any age	May be family	Joint contractures,	Cardiac complications	Cardiac echo, ECG +/- Holter, FVC in sitting and lying.
		history of	rigid spine	probably invariable:	High risk of ventricular arryhythmia after pacing so
		sudden death	The state of the s	arrhythmia, subsequent	implantable defibrillator indicated. Medical management of
				cardiomyopathy,	cardiomyopathy, respiratory support
				respiratory failure	
LGMD1C	Any age	Myalgia, muscle	PIRCs, rippling,	Not reported with	
		rippling	muscle	LGMD1C phenotype	

			hypertrophy		
LGMD2A	Typically	May present	Scapular winging,	Not frequent	Q
	2-15: in	with toe	some patients		
	some	walking	have contractures,		O'
	patients		often atrophic		
	older				
LGMD2B	Most	Inability to walk	Usually calf	Not frequent	
	present	on tip-toe.	wasting, biceps		
	between	Around 10%	wasting. Shoulder	,47	
	17-25	may present	girdle weakness		
	years	with calf pain	usually later than		
		and swelling	proximal/ distal)	
			lower limb		
			weakness		
LGMD2C-F	May	May present in	May be muscle	Cardiomyopathy and	Cardiac echo and ECG on annual basis, FVC particularly
	present	Duchenne/	hypertrophy,	respiratory failure after	after loss of ambulation, pulse oximetry. Medical
	any age:	Becker like way	scapular winging	confinement to	management of cardio-myopathy, nocturnal respiratory

	most in			wheelchair	support
	child-				Ŕ
	hood				2
LGMD2G	2 nd		May be distal	Not reported	0
	decade		weakness		
LGMD2H	2 nd			Not reported	
	decade				
LGMD2I	Wide	Clinically may	Muscle	Cardiomyopathy and	Cardiac echo and ECG on annual basis, FVC in sitting and
	range of	resemble	hypertrophy	respiratory failure	lying at all ages, pulse oximetry. Medical management of
	severity	Duchenne/		(diaphragmatic) possible	cardio-myopathy, nocturnal respiratory support
		Becker MD		even while still ambulant	
LGMD2J	1 st	History of distal		Late respiratory	
	decade	weakness in	8	complications	
		heterozygotes	*		
LGMD2K	1 st	May present	May be mental	Full spectrum of	
	decade	early with	retardation,	complications possibly	
		global delay	microcephaly.	not yet appreciated	

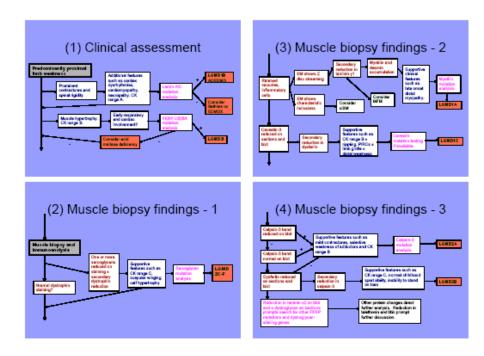
Upper li	mb	4
weaknes	ss may be	, Q
worse th	an lower	2
limb we	akness	,C`

Table 3. Diagnostic features: creatine kinase and muscle biopsy findings.

Disease	Creatine kinase	Histology	Primary abnormality on	Secondary abnormality on
	levels		immunolabelling	immunolabelling
LGMD1A	Usually normal or	Dystrophic, may be	myotilin N	↓ laminin γ1
	mild elevation	desmin accumulation,		
		rimmed vacuoles	.65	
LGMD1B	Typically normal or	dystrophic	lamin A/C N	↓ laminin β1
	mild elevation:			
	severe cases may be			
	higher			
LGMD1C	From 3- 10x normal	myopathic, dystrophic	↓ caveolin-3	↓ dysferlin
		141		
LGMD2A	Often at least 10x	dystrophic	Variable loss on	
	normal		immunoblotting	
LGMD2B	May be up to 100x	Dystrophic, may be	↓ dysferlin	(may be↓ calpain 3, and/or
	normal	inflammatory features		↓ caveolin 3)
LGMD2C-	Often at least 10x	dystrophic	↓ sarcoglycans	↓ other SGs, dystrophin
F	normal			
LGMD2I	Often at least 10x	dystrophic	FKRP antibody not in	↓ laminin α2, αDG
	normal		routine use	
LGMD2J	Few cases: high CK	Myopathic, dystrophic,	↓ titin	↓ calpain-3
		may be rimmed		
		vacuoles		
LGMD2K	At least 10x normal	dystrophic	POMT1 antibody not in	↓ laminin α2, αDG
			routine use	



Figure





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