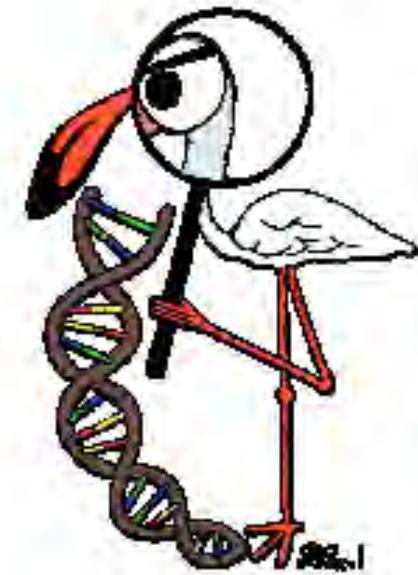


Studio multicentrico sulle Distrofie Muscolari dei Cingoli Telethon-UILDM

Centro Dino Ferrari, Fondazione IRCCS Ca' Granda
Ospedale Maggiore Policlinico, DEPT, UO
Neurologia, Università di Milano



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Ente Sanitario Regionale

Regione Lombardia



UNIVERSITÀ
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DI MILANO

BRAIN

VOL. 77, PART 2.

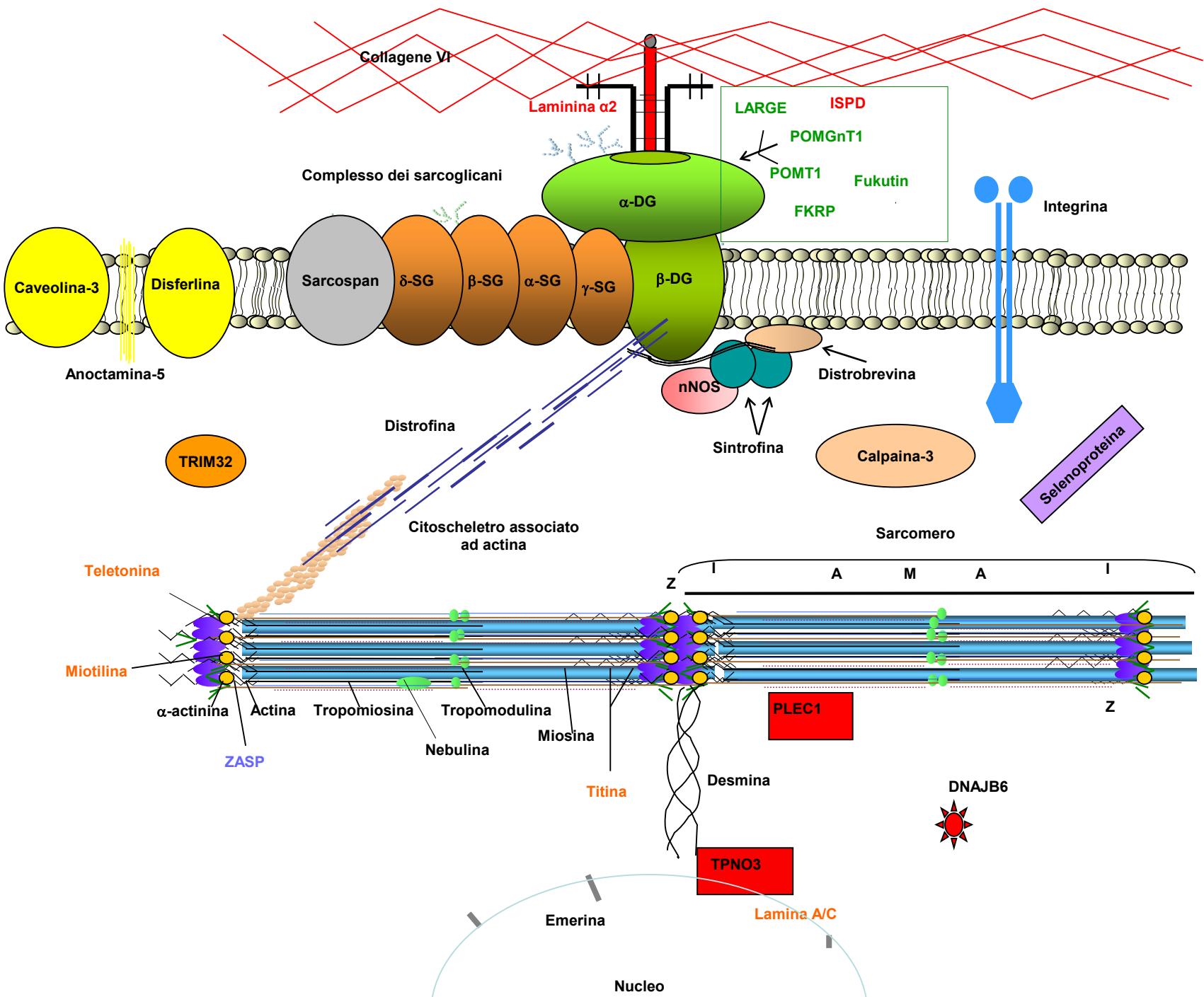
ON THE CLASSIFICATION, NATURAL HISTORY
AND TREATMENT OF THE MYOPATHIES

BY

JOHN N. WALTON AND F. J. NATTRASS

(*From the Department of Medicine, King's College, University of Durham and the
Royal Victoria Infirmary, Newcastle upon Tyne*)

In our opinion, these cases should be regarded as a distinct clinical and genetic group and we suggest that they should be called *limb-girdle muscular dystrophy*. Leaving aside certain clinical characteristics which may be added later, we feel that the cardinal features of this type are (a) onset usually late in the first or in the second or third decade but sometimes in middle age, (b) commencement of muscular weakness in either the shoulder or pelvic girdle, (c) transmission usually via an autosomal recessive gene, (d) a relatively slow course which nevertheless leads to severe disablement and often death before the normal age.



Molecular classification

	Gene	Gene location	Protein
LGMD 1A	MYOT	5q31	Myotilin
LGMD 1B	LMNA	1q21.2	Lamin A/C
LGMD 1C	CAV3	3p25	Caveolin-3
LGMD1D	DES	2q35	Desmin
LGMD 1E	DNAJB6	7q36	DnaJ homolog subfamily B member 6
LGMD 1F	TNPO3	7q32	Transportin3
LGMD 1G	HNRPDL	4p21	Heterogeneous nuclear ribonucleoprotein D-like protein
LGMD 1H	?	3p23-p25	?

- 8 Autosomal Dominant (LGMD1)
- 23 Autosomal Recessive (LGMD2)



30% without a molecular diagnosis

	Gene	Gene location	Protein
LGMD 2A	CAPN3	15q15.1-q21.1	Calpain-3
LGMD 2B	DYSF	2p13.3-913.1	Dysferlin
LGMD 2C	SGCG	13q12	γ-Sarcoglycan
LGMD 2D	SGCA	17q12-q21,33	α-Sarcoglycan
LGMD 2E	SGCB	4q12	β-Sarcoglycan
LGMD 2F	SGCD	5q33	δ-Sarcoglycan
LGMD 2G	TCAP	17q12	Telethonin
LGMD 2H	TRIM32	9q31-9q34	TRIM32
LGMD 2I	FKRP	19q13,3	Fukutin Related Protein
LGMD 2J	TTN	2q24.3	Titin
LGMD 2K	POMT1	9q34.1	O-Mannosyl transferase-1
LGMD 2L	ANO5	11p12-p13	Anoctamin-5
LGMD 2M	FKNT	9q31-q99	Fukutin
LGMD 2N	POMT2	14q24	O-Mannosyl transferase-2
LGMD 2O	POMGnT1	1p34	Protein O-mannose beta-1,2-N-acetylglucosaminyl-transferase
LGMD 2P	DAG1	3p21	Dystrophin-associated glycoprotein
LGMD 2Q	PLEC1	8q24.3	Plectin
LGMD 2R	LAMA2	6q22	Merosin
LGMD2S	TRAPPC11	4q35	Transport protein particle complex 11
LGMD2T	GMPPB	3p21	GDP-mannose pyrophosphorylase B
LGMD2U	ISPD	7p21	Isoprenoid synthase domain containing protein
LGMD2V	POMK	8p11.21	Protein O-mannose kinase
LGMD2W	PINCH2/LIMS2	2q14.3	Particularly interesting new cys-his protein 2

Clinical and laboratory network for LGMD diagnosis, in view of a national registry Telethon GUP10006



- Department of Neurological Sciences, I.R.C.C.S. Foundation Cà Granda Ospedale Maggiore Policlinico, University of **Milan**
- NeuroMuscular Unit-IRCCS E Medea **Bosisio Parini**,
- Department of General Pathology, University of Naples, **Naples**
- Telethon Institute of Genetics and Medicine, **Naples**
- IRCCS Fondazione "San Camillo" Hospital, Lido di **Venezia**
- Department of Neurosciences, University of **Torino**
- Neuromuscular Diseases and Neuroimmunology Unit, Fondazione IRCCS Istituto Neurologico C. Besta, **Milan**
- Department of Neurosciences, Psychiatry and Anaesthesiology, University of Messina, **Messina**
- Department of Neurological Sciences, **Verona**
- Department of Clinical and Experimental Medicine, University of **Pisa**
- Center of Myology and Neurodegenerative Diseases, Istituto Giannina Gaslini, **Genova**
- Department of Neurology, Policlinico Universitario A. Gemelli, University Cattolica del Sacro Cuore of **Rome**
- Department of Neurosciences, University of **Padua**
- IRCCS Fondazione Stella Maris, Calambrone, **Pisa**

Objectives

- Create a national registry of LGMD Italian patients
- Collect clinical data about neuromuscular, respiratory, cardiac, cognitive involvement
- Delineate natural history of each sub-group
- Define outcome measures
- Further investigate patients without molecular diagnosis

Patient Selection

INCLUSION CRITERIA:

1. limb girdle and proximal upper and lower limb muscle weakness
(Proximo-distal phenotypes may be included)
2. dystrophic pattern at muscle biopsy and/or
3. proven deficiency of LGMD proteins at IHC or WB analysis and/or molecular confirmation

EXCLUSION CRITERIA:

Limb Girdle syndrome with identified alternative etiology (inflammatory, metabolic myopathies, congenital myopathies, congenital myasthenic syndromes, SMA type II and III, FSH, dystrophinopathies, DM1, DM2 and mitochondrial myopathies)

Data collection

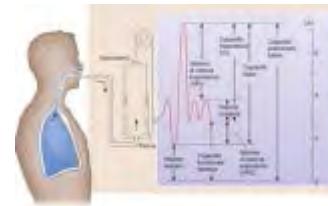
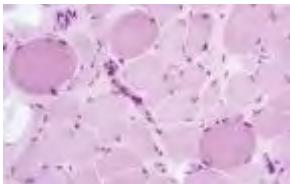


ANAMNESTIC DATA

Familiarity
Onset
CK values
Muscular involvement
(distribution)
Tendon retraction
Scoliosis

MUSCLE BIOPSY

Morphology
IHC analysis
WB analysis



RESPIRATORY FUNCTION

Spirometry
Nocturnal saturimetry
NIV



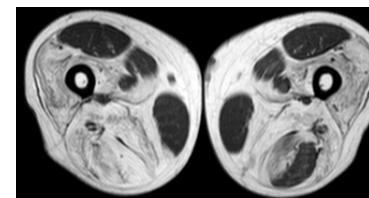
FUNCTIONAL EVALUATIONS

MRC
Walton
MFM
6MWT



MOLECULAR ANALYSIS

CARDIAC FUNCTION
ECG
Holter ECG
Echocardiogram



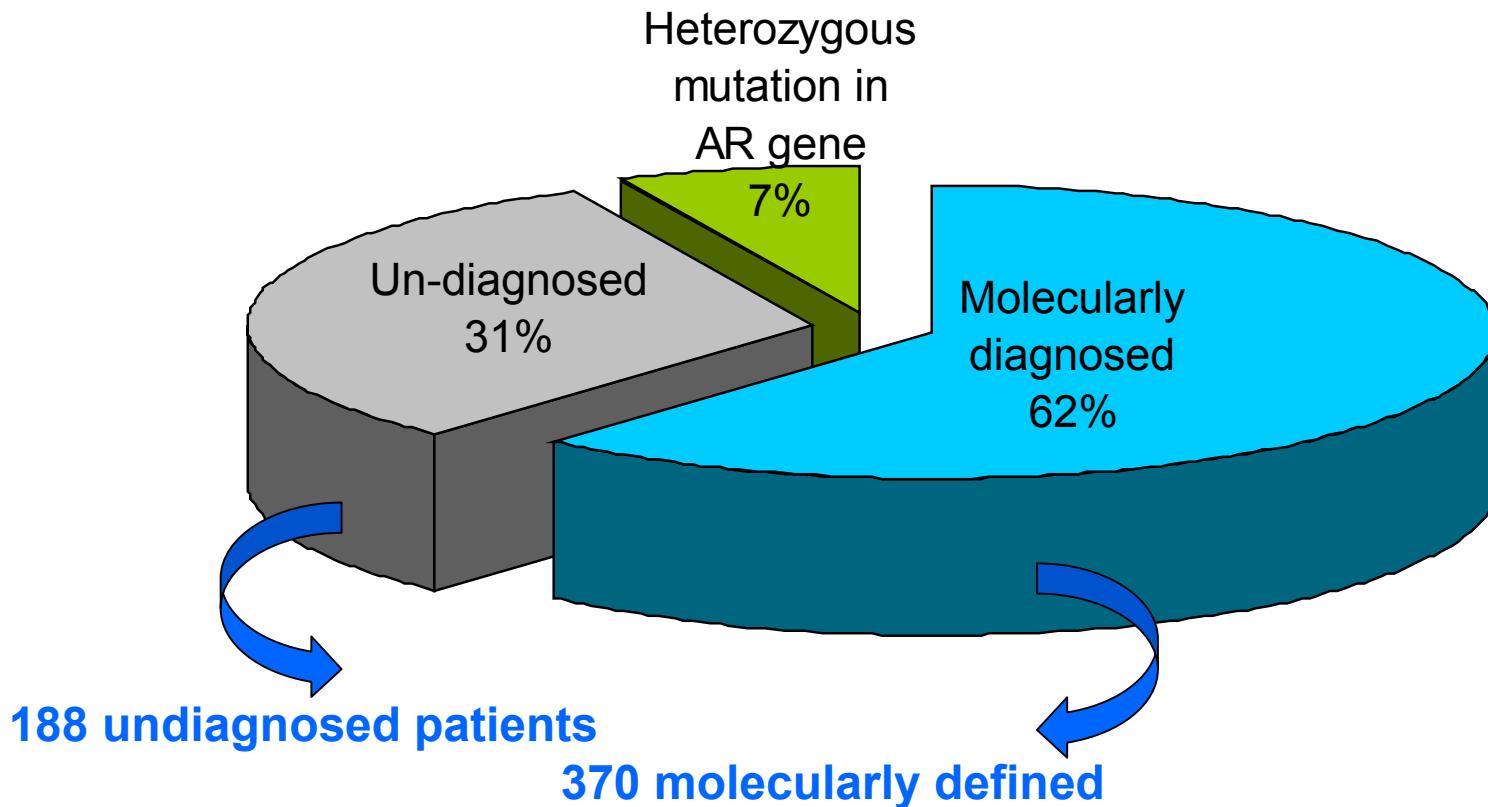
MAGNETIC RESONANCE

Muscular MRI
Brain MRI

DATABASE containing retrospective and prospective data

LGMD National registry

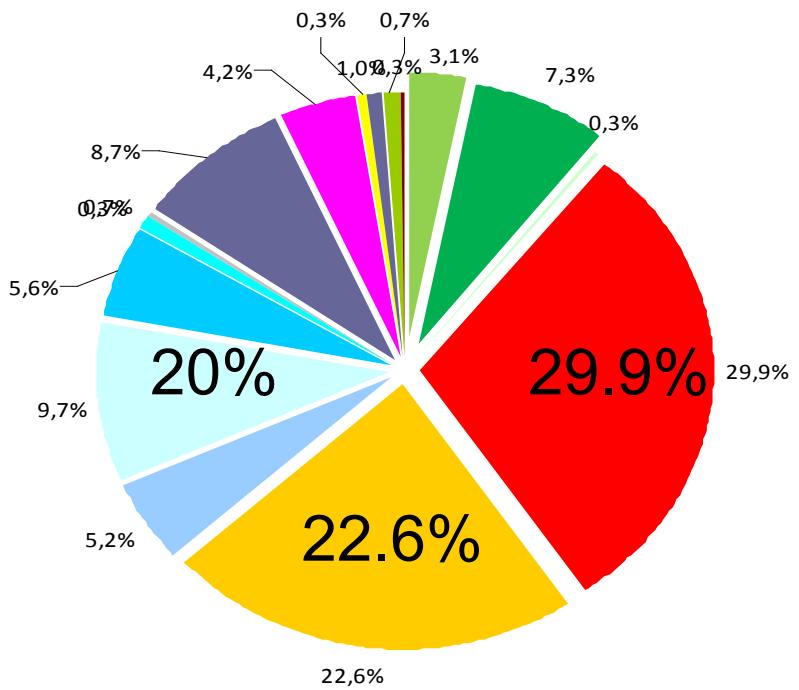
599 LGMD patients



LGMD registry

370 molecularly defined

Probands



LGMD1B

LGMD1C

LGMD1D

LGMD2A

LGMD2B

LGMD2C

LGMD2D

LGMD2E

LGMD2F

LGMD2G

LGMD2I

LGMD2L

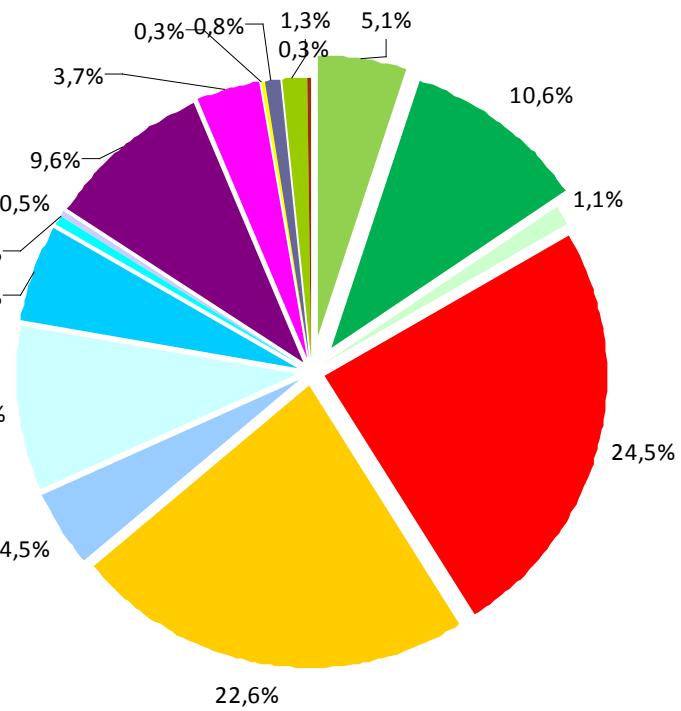
LGMD2J

LGMD2 POMT2

LGMD2 LAMA2

LGMD2 ISPD

Patients

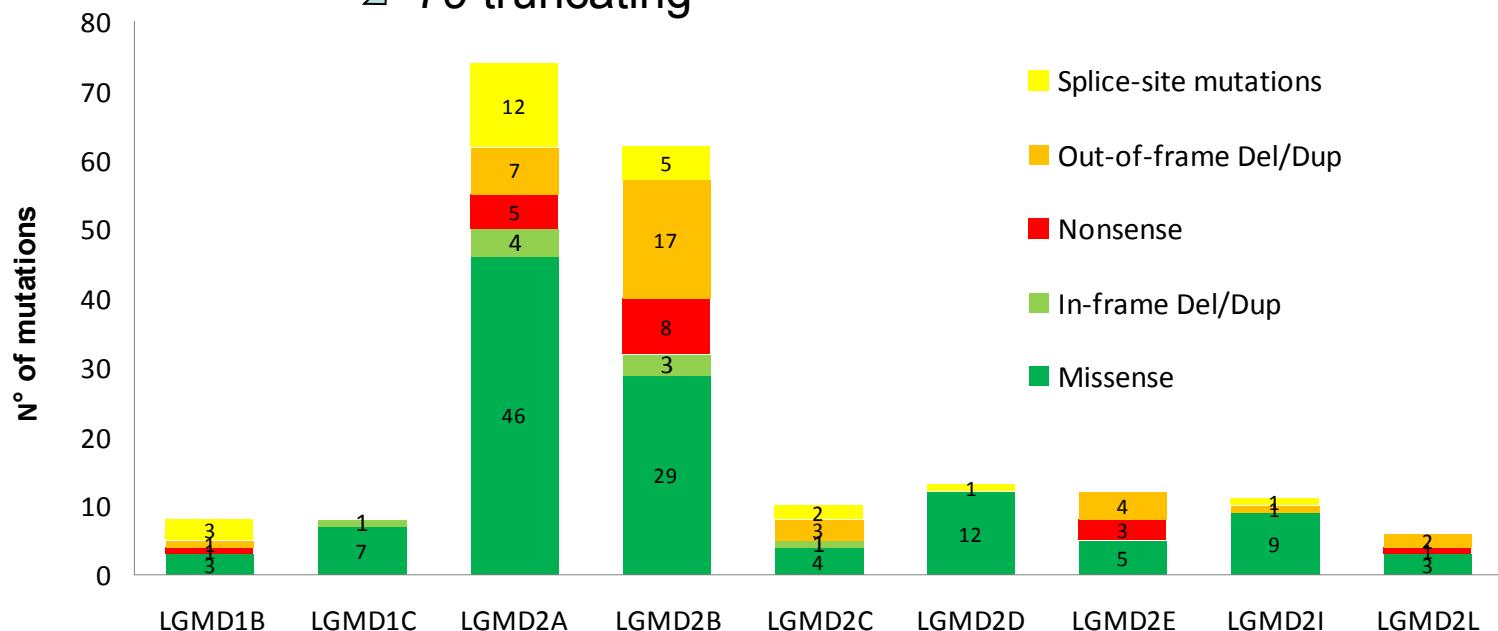


Mean follow-up $17,0 \pm 11,6$ years

Molecular aspects

207 different mutations → 128 in-frame

→ 79 truncating



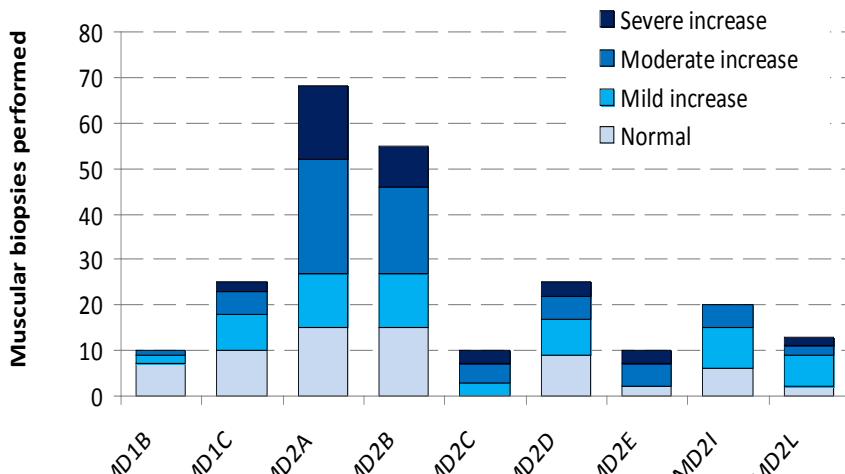
No hot-spots

Most frequent mutations:

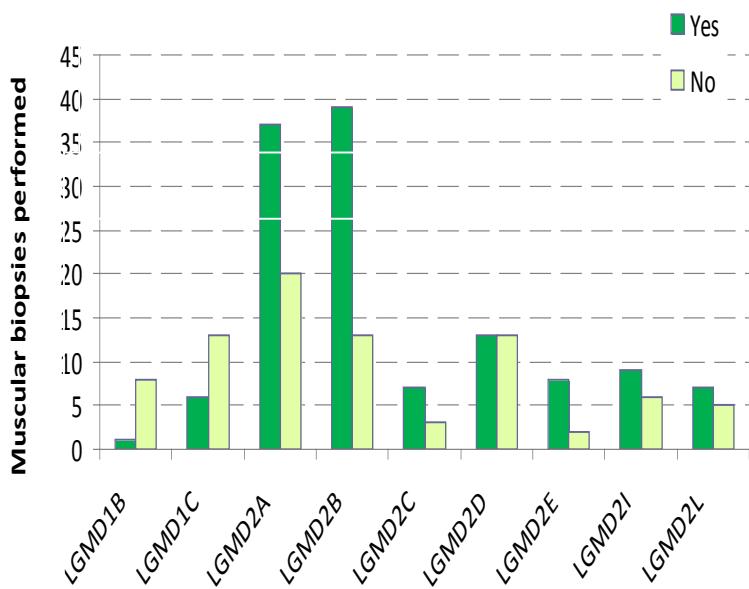
- c.826C>A (p.Leu276Ile) exon 4 (*FKRP*) → 23/38 alleles (60%) LGMD2I
- c.525delT (p. Phe175LeufsX20) exon 6 (*SGCG*) → 8/22 (36%) LGMD2C
- c.850C>T (p.Arg284Cys) exon 7 (*SGCA*) → 16/50 (32%) LGMD2D
- c.377_384dup (p.Gly128GlnfsX2) exon 3 (*SGCB*) → 9/30 (30%) LGMD2E

Muscle biopsy

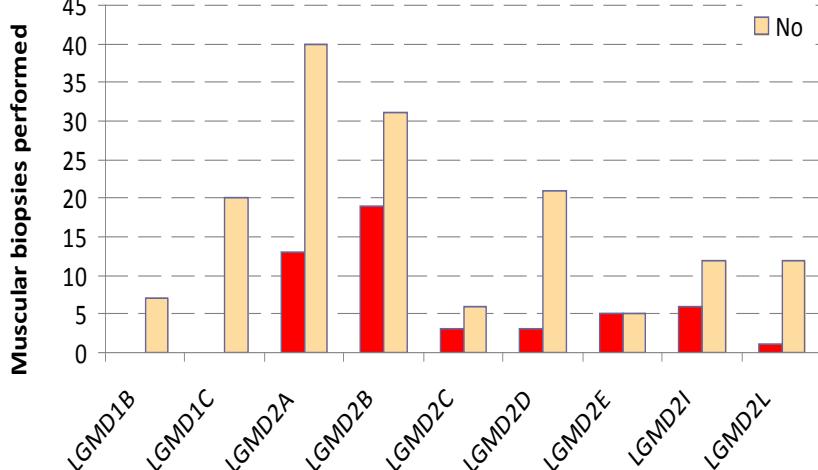
Connective Tissue



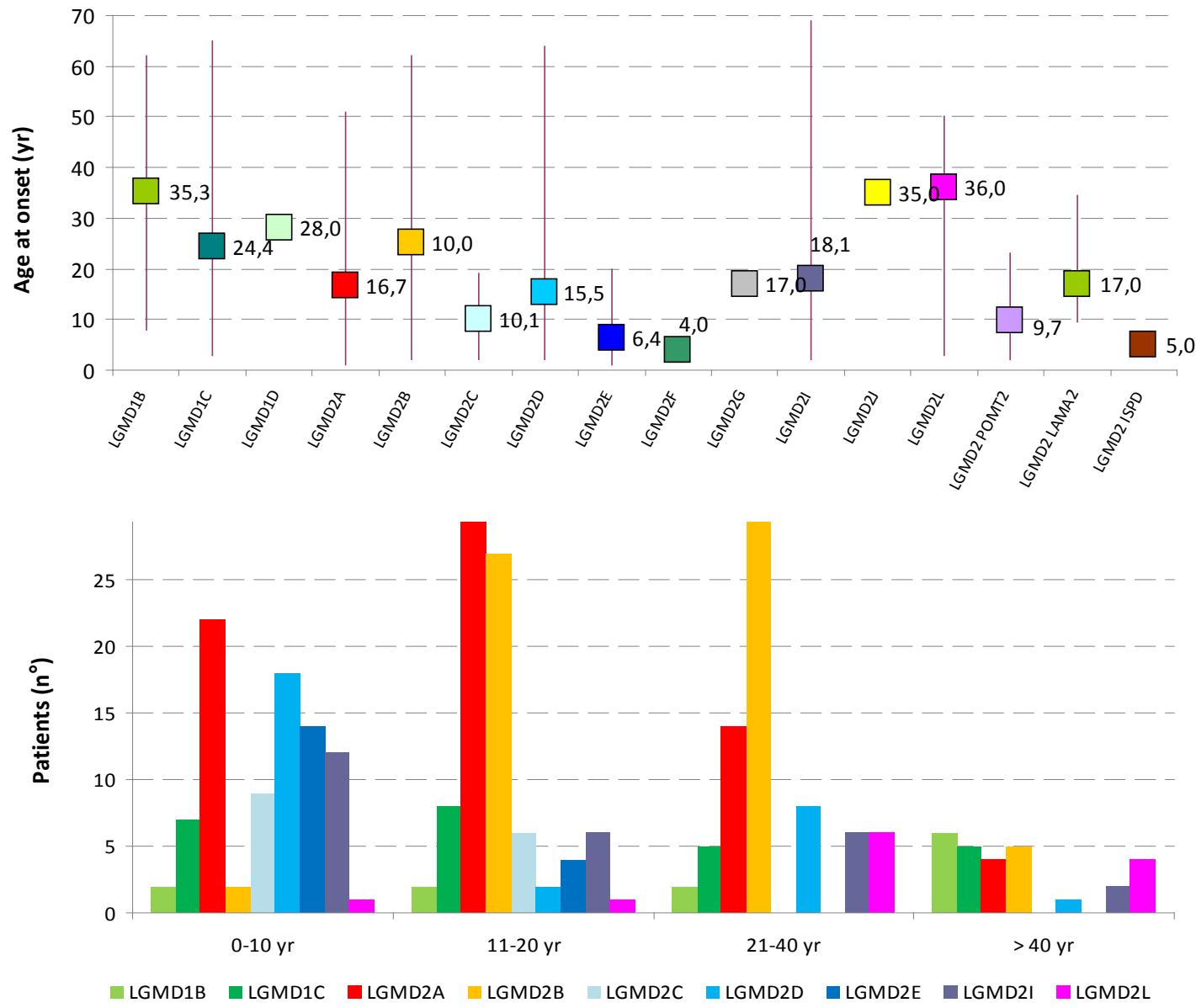
Necrosis



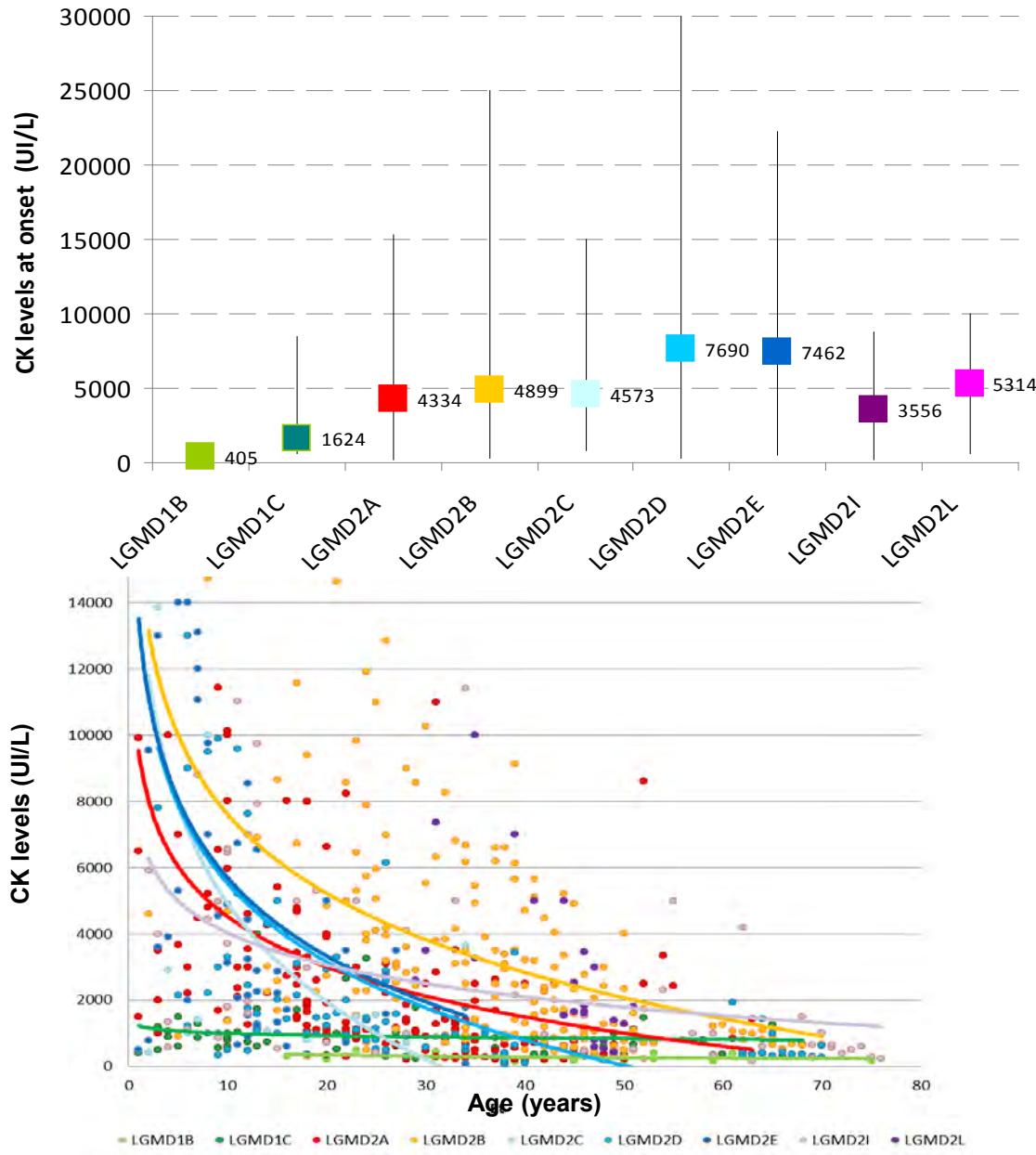
Inflammation



Clinical evolution – Age of onset

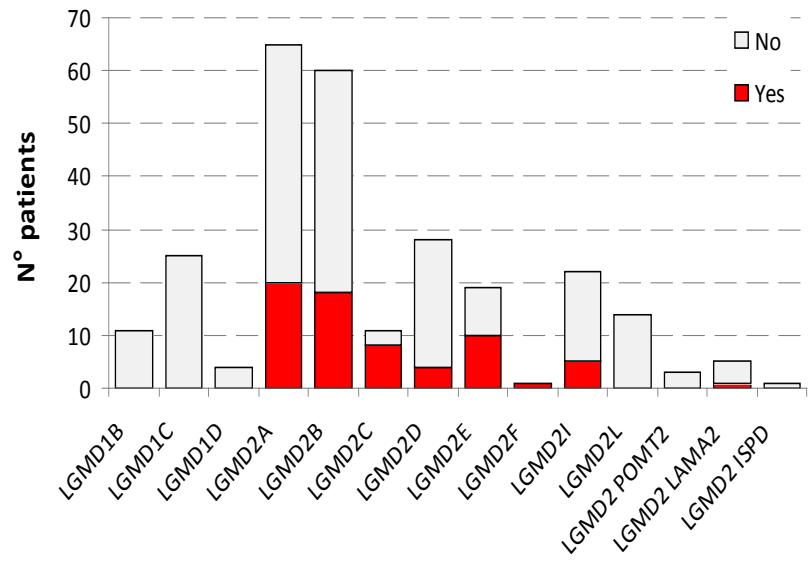


Clinical evolution – CK levels

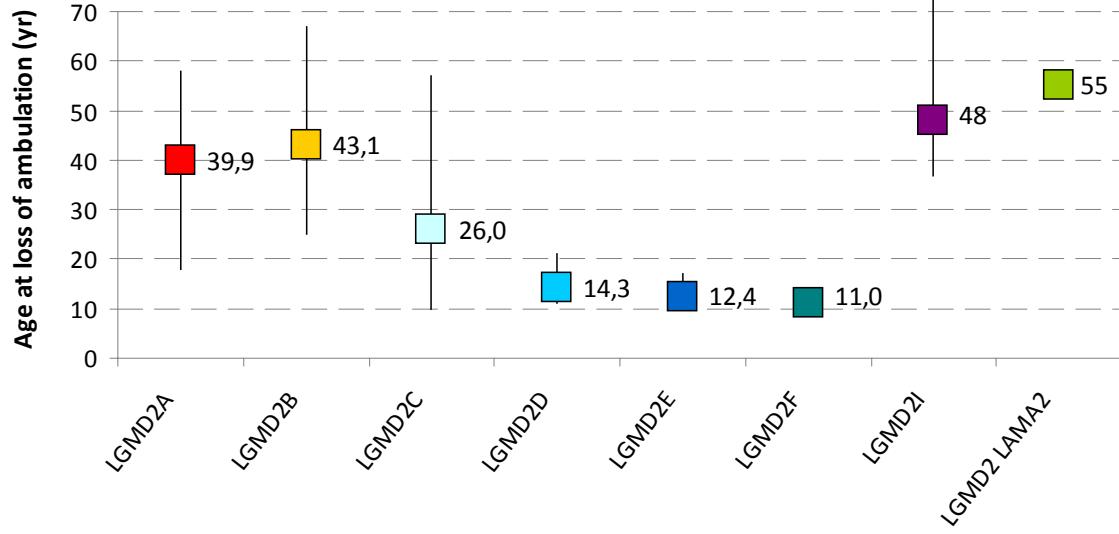
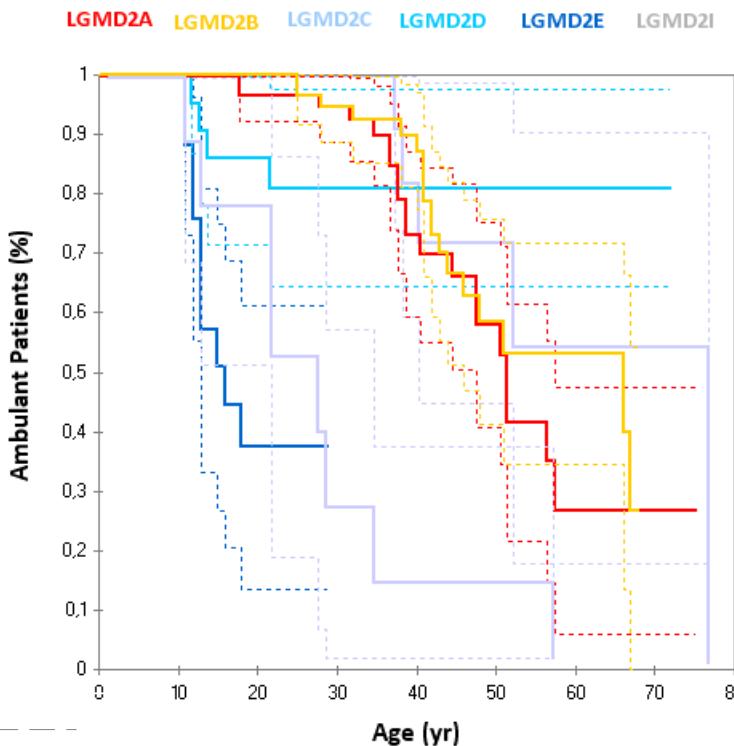


Clinical evolution – Loss of independent ambulation

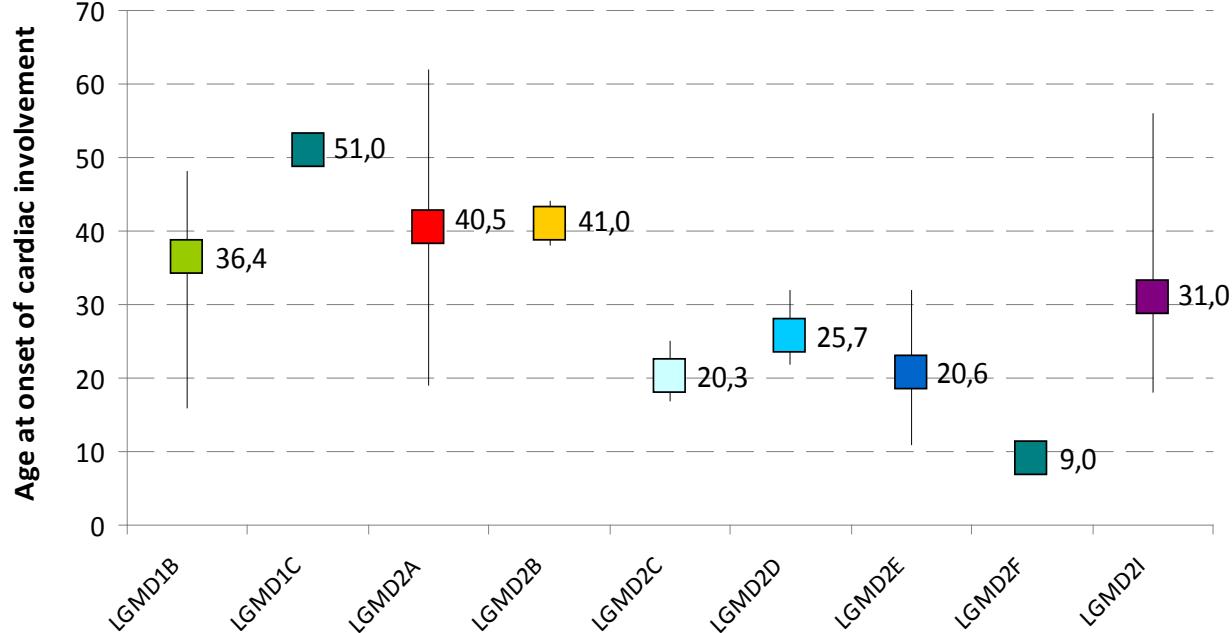
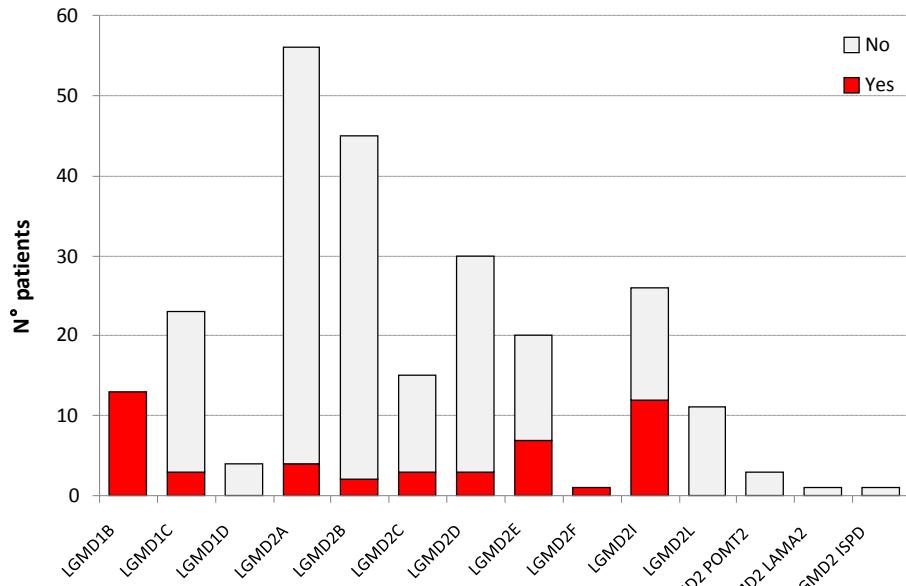
Loss of ambulation



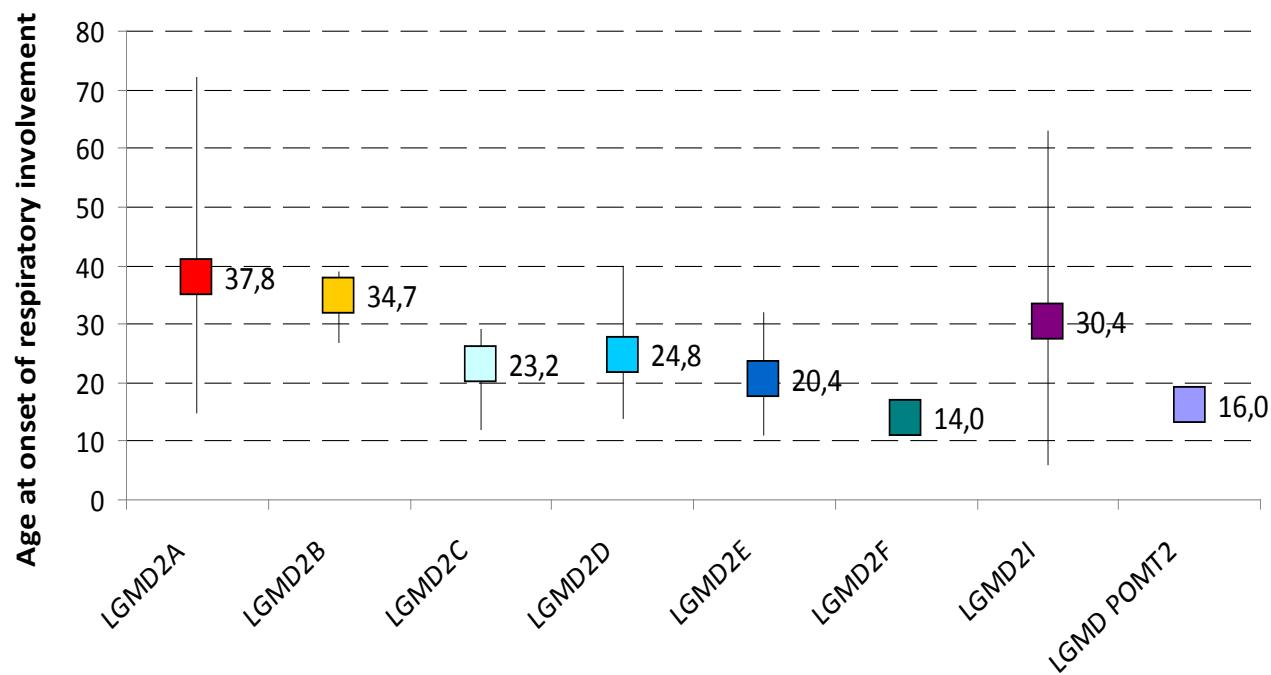
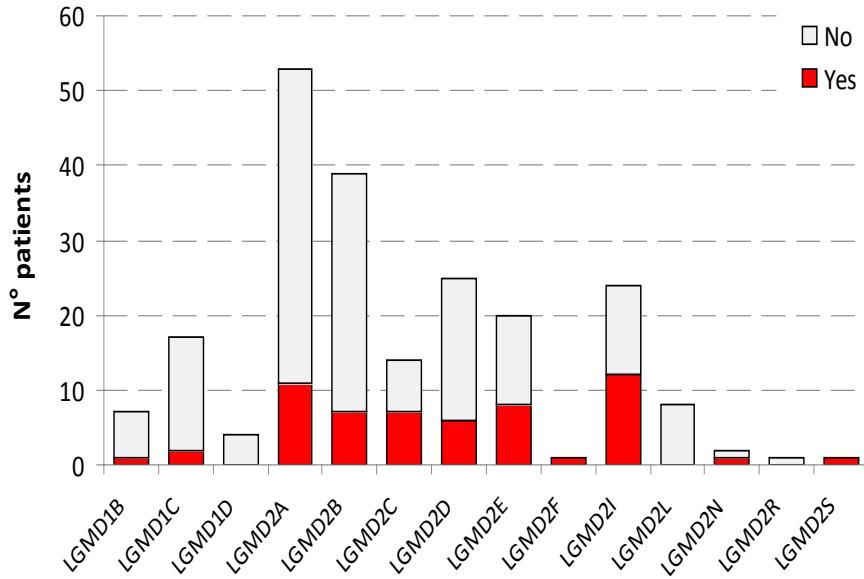
C



Clinical evolution – Cardiac involvement



Clinical evolution – Respiratory involvement



LGMD differential diagnosis

Age at onset

- Earlier onset in LGMD2C-2E, dystroglycanopathies
- Later onset LGMD2I and 2L

CK levels

- High levels in LGMD2B, LGMD2I
- Low levels in LGMD1B, 1C

Ambulation loss

- Early in LGMD2D, 2E
- Late LGMD2A, 2B, 2C

Cardiac involvement

- LGMD2E , LGMD2C, LGMD2F, LGMD2I, LGMD2M

Respiratory involvement

- LGMD2A, LGMD2D, LGMD2M

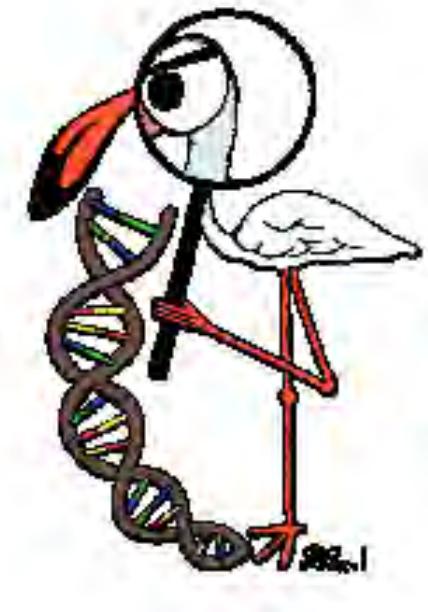
Muscle biopsy analysis

- LGMD1C, 2A, 2B, sarcoglycanopathies, dystroglycanopathies

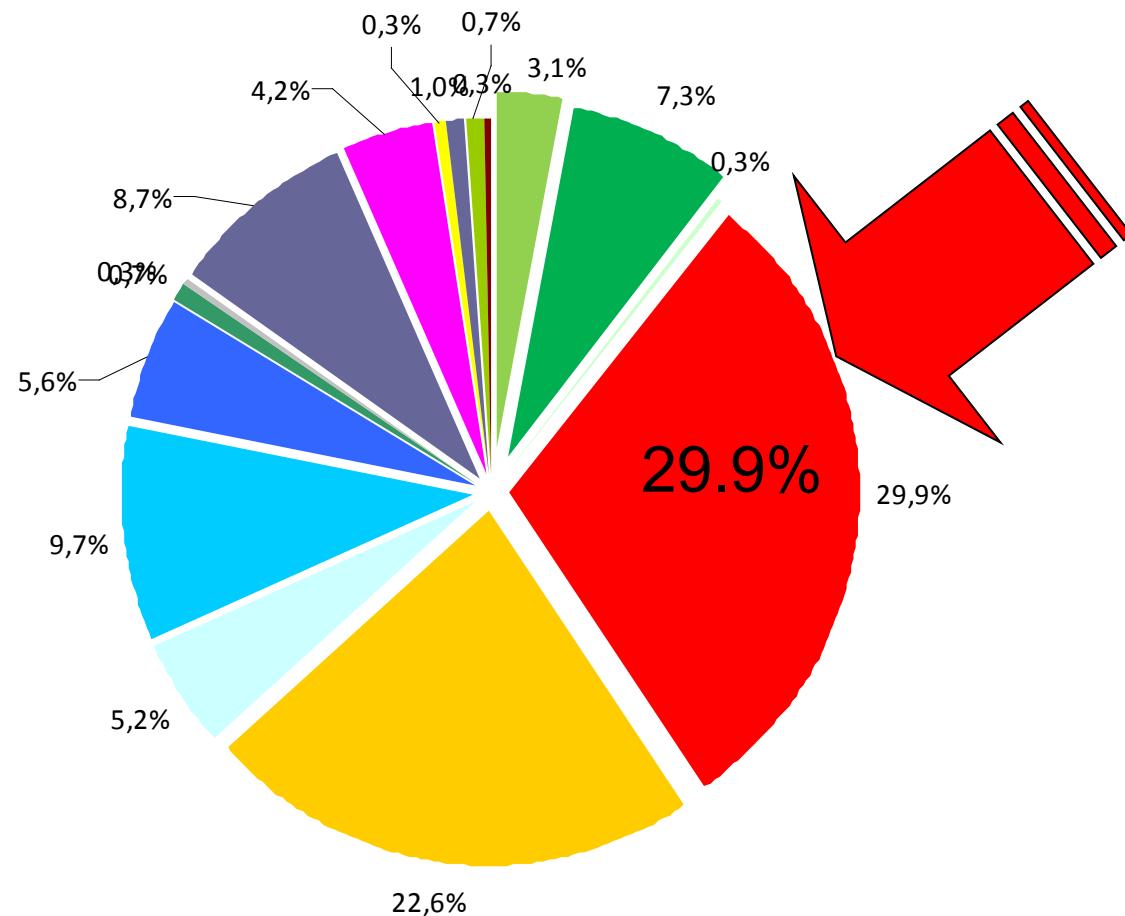
LGMD national registry

- LGMD molecular epidemiology in Italy is extremely heterogenous.
- The relative frequency of the predominant LGMD types differs in Southern Europe from that other Northern countries
- Decreasing the proportion of genetically undiagnosed cases is probably possible by Next Generation Sequencing.
- New common or rare pathogenetic pathways may emerge from this effort.
- Building an itemised disease natural history database for at least 30 different diseases (often with allelic early onset and late onset phenotypes) is a complex and time consuming task.
- Therapeutic trials involving this population are scarce and reflect into less attention paid to the data collection to establish a reference clinical database for these disorders.

What about LGMD2A?



LGMD2A POPULATION



86 probands
92 patients

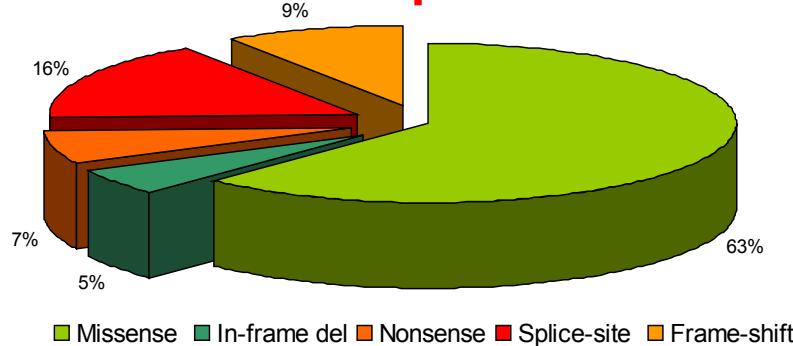
F:M= 1.3/1

Mean follow-up
 18.8 ± 9.2 years

Age of last evaluation
 37.1 ± 14.2 years

LGMD2A Diagnostic clues

Molecular aspects



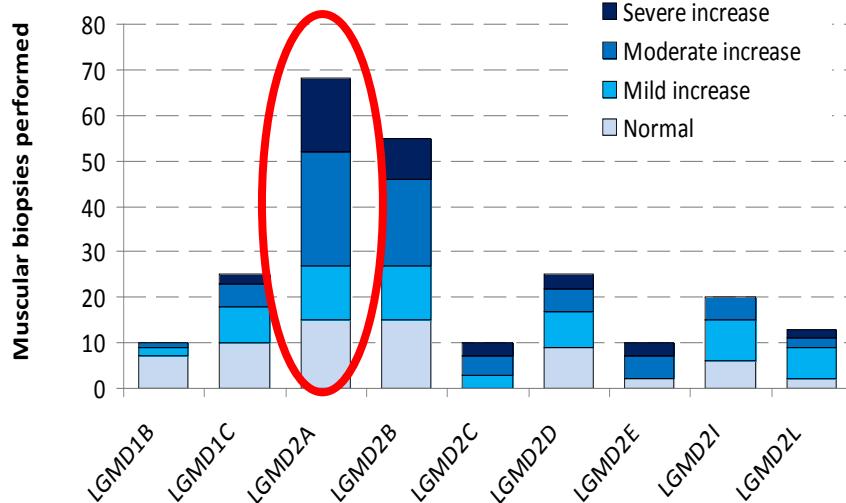
69% of mutations are located in exons 4,11,10,13,1,21,5,6

Most frequent mutations:

c.550delA	6.8%
c.1469G>A	6.1%
c.2242C>T	4.5%

Biooptical aspects

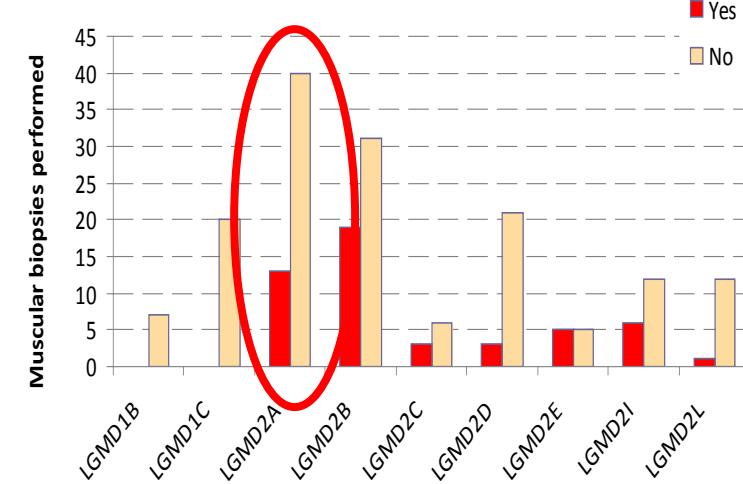
Connective Tissue



10 patients without connective tissue increase
(7 quadriceps femori)

No correlation with age or disease duration

Inflammation

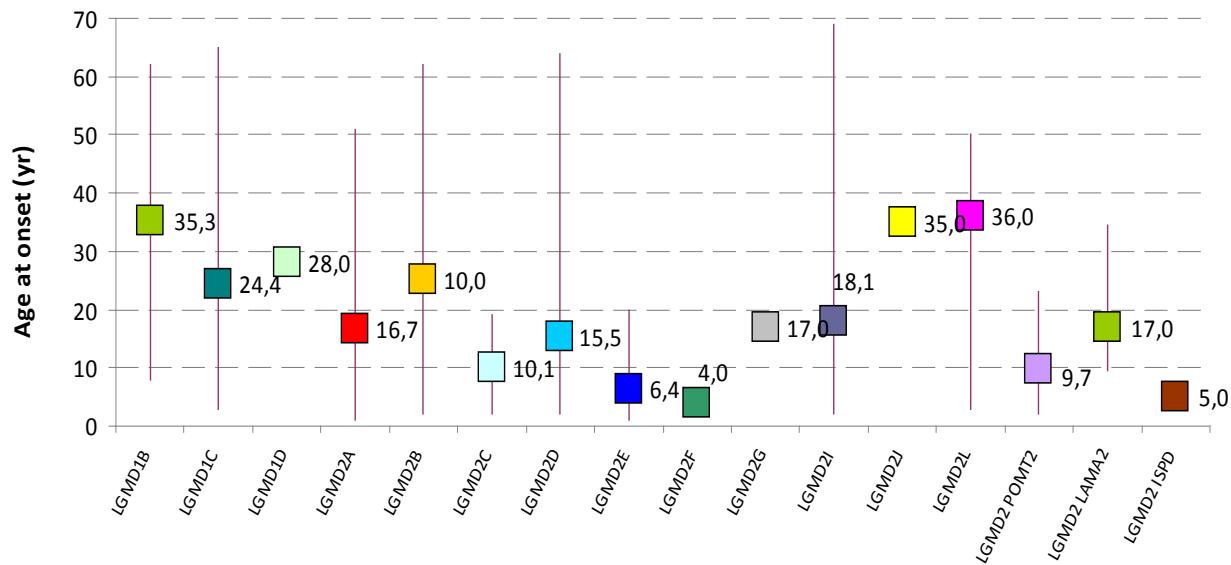


Protein analysis

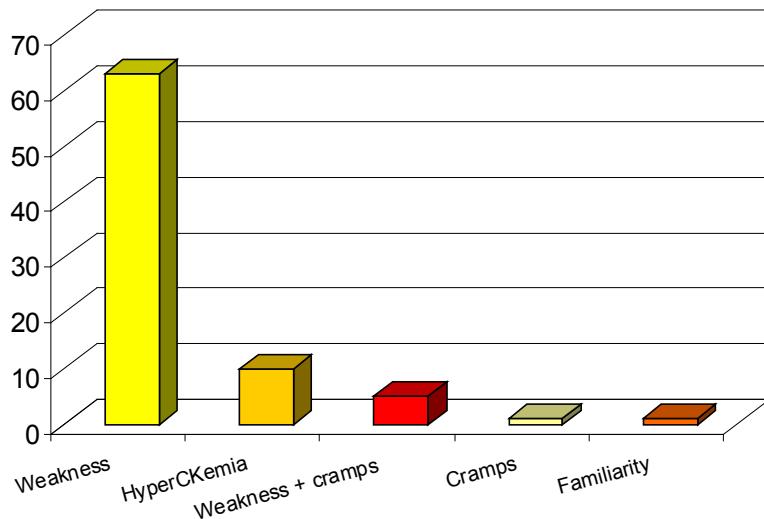
Western-blot analysis: reduction in 57 patients,
normal expression in 3 subjects

LGMD2A Clinical evolution 1

Age at onset



Symptoms at onset



Onset of weakness

17.6 ± 11.8 years
(range 1-51).

< 1 years of age
≤ 20 years of age
20-40 years of age
> 40 years of age

→3 pt (5%)
→47 pt (75%)
→12 pt (19%)
→4 pt (6%)

LGMD2A Clinical evolution 2

Distribution of muscular involvement

55 patients pelvic + shoulder girdle

8 patients pelvic girdle (4-75 years old, 3-30 years from onset)

3 patient early distal involvement

Difficulties in postural changes

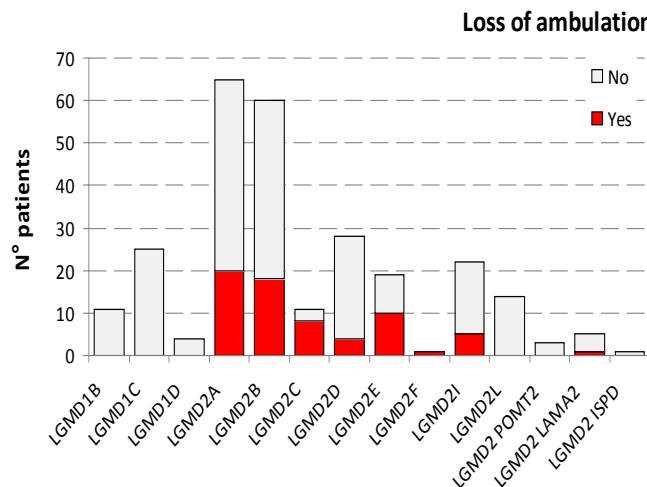
27/42 → 31.9 ± 15.9 years

17.2 ± 7.0 years from onset

Loss of ambulation

20/65 → 31% mean age 39.9 ± 11.5 years

Wheelchair-bound patients have earlier onset (15.4 ± 8.3 vs 19.6 ± 12.1 years)



LGMD2A Clinical evolution 3

CK values

$4025 \pm 3393 \text{ UI/L}$ (range 217-15300)

Cardiac involvement

4/56 → 7%

Mean age at onset 39 ± 22 years

Onset respectively at 19, 20, 55 and 62 years of age

Range 8-17 years from disease onset

2 patients with dilatative cardiomyopathy

Furthermore 12 patients with rhythm alteration

Respiratory involvement

11/53 → 21%

Mean age at onset 37.8 ± 22.8 years

Restrictive pattern

Peculiar aspects

Tendon retraction 29/51 64%

Sural hypertrophy 15/49 pt

Scoliosis 16/44

Thank you



UO Neurologia - IRCCS Ca' Granda–Milano

Prof. Bresolin, Prof. Comi, Dr. Magri, Prof. Corti, Dr. Govoni, Dr. Brusa, Dr. Del Bo, Dr. Ronchi, Dr Piga

UOD Neuromuscolari, IRCCS Ca' Granda, University Milan

Prof Moggio, Dr. Sciacco, Dr. Colombo, Dr Peverelli, Dr Villa

Ist. E. Medea Bosisio Parini – Lecco

Dr. D'Angelo, Dr. Gandossini

Dr. Brighina, Dr. Micoli

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Telethon Institute of Genetics and Medicine (TIGEM) – Napoli

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Dr. Tomelleri, Dr Tonin*

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Dr. Fiorillo*