# Gamma-sarcoglycanopathy (LGMDR5): clinical and genetic study of a pediatric Tunisian cohort

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## INTRODUCTION

•Limb-girdle muscular dystrophy **R5** type (LGMDR5) is an autosomal-recessive disorder caused by mutations in gamma-sarcoglycan encoding gene (1,2)

•It is characterized by childhood onset of progressive muscular dystrophy (1)

•It is **frequent in North African** populations (1,2)

#### **OBJECTIVES**

clinical determine and genetic the characteristics of a Tunisian series of LGMDR5

## MATERIALS AND METHODS

2022 •Retrospective study from 2004 to collecting demographic, clinical, biological, electrophysiological, and genetic data of patients with genetically confirmed LGMDR5

#### RESULTS

- •24 patients
- •Demographic and clinical features (Table 1)
- •Explorations (Table 2)
- •**Treatment**: Steroids at the dose of 0.75mg/kg/d
- : 10d ON/10d OFF (83% of patients)
- •Mean **follow-up** period: 3 years
- •Evolution (figure 1)

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#### Table 2: Result of explorations

Explo

Serum

Transa Electro Muscle

Geneti

1-El Kerch F, Ratbi I, Sbiti A, et al. Carrier frequency of the c.525delT mutation in the SGCG gene and estimated prevalence of limb girdle muscular dystrophy type 2C among the Moroccan population. Genet Test Mol Biomarkers. 2014 Apr;18(4):253-6 2- Kefi M, Amouri R, Driss A, et al. Phenotype and sarcoglycan expression in Tunisian LGMD 2C patients sharing the same del521-T mutation. Neuromuscul Disord. 2003 Dec;13(10):779-87

#### Table 1: Demographic and clinical features

graphic and clinical features		N° of patients/fam
patients		24
families		18
nale/female		11/13
nguinity		17 families
al history of similar case		8 families
age of onset (years)		3.7
age at last evaluation (years)		9,9
iral on	Difficulties to run and to climb stairs	18
	Gait disturbance	4
	High CK level	2
nation	Proximal lower limb weakness	24
	Calf hypertrophy	20
	Cognitive impairment	3

ration	Result (% patients)
creatine kinase levels	Levels ranged from 2280 to 18461 U/L. (NV (100%)
minase	High level of AST and ALT (100%)
myography	Myogenic pattern (100%)
e biopsy	Dystrophic changes (17%)
c study	Homozygous c.521delT mutation in the SG gene (100%)

#### REFERENCES





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