



ALAG

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DE OCTUBRE 2019

MENDOZA, ARGENTINA

La arquitectura
del genoma:
su expresión en
los fenotipos
y las poblaciones

XVII Congreso Latinoamericano de Genética
XLVII Congreso Argentino de Genética
LII Reunión Anual de la Sociedad de Genética de Chile
VI Congreso de la Sociedad Uruguaya de Genética
V Congreso Latinoamericano de Genética Humana y
V Simposio Latinoamericano de Citogenética y Evolución

CLUSTERS OF MPS IN LATIN AMERICA

FRANCYNE KUBASKI, MSC, PHD

DISCLOSURES

- I have been in receipt of honoraria for lectures on LSDs from Sanofi-Genzyme Corp
- Lucid and Biomarin consultant
- Informed consent was obtained for image use

OUTLINE

- MPS introduction
- GAGs
- MPS management
- Clusters of MPS in Latin America

MUCOPOLYSACCHARIDOSES

- Mucopolysaccharidoses (MPS) are rare lysosomal storage disorders caused by deficiency of lysosomal hydrolases leading to the accumulation of undegraded glycosaminoglycans (GAGs) (Table I). With the exception of MPS II, which is X-linked, the MPS are autosomal recessive.

Table I – Classification of the MPS approached in this study

Number	Enzyme deficiency	Eponym	Storage material	Gene
IIIB	α -N-acetylglucosaminidase	Sanfilippo B	Heparan sulfate	<i>NAGLU</i>
IIIC	Acetyl-coA- α -glucosaminide acetyltransferase	Sanfilippo C	Heparan sulfate	<i>HGSNAT</i>
IVA	N-acetylgalactosamine-6-sulfate sulfatase	Morquio A	Chondroitin-6-sulfate & Keratan sulfate	<i>GALNS</i>
VI	N-acetylgalactosamine-4-sulfatase	Maroteaux-Lamy	Dermatan sulfate	<i>ARSB</i>

GLYCOSAMINOGLYCANS (GAGS) ARE IMPORTANT FOR CELLULAR FUNCTION

- Gags are important structural components of cellular membranes, extracellular matrix, connective tissue, and joint fluids
- At the cellular level, GAGs are involved in cell division, differentiation, and cell-cell communication
- In healthy individuals, GAGs are degraded by lysosomal enzymes in a stepwise fashion within the lysosome



GLYCOSAMINOGLYCANS- GAGS

Category	Uronic acid	Hexosamine	Modifications
Heparin/heparan	Iduronic acid/glucuronic acid	Glucosamine	X-sulfated Y-acetylated/sulfated
Chondroitin/ dermatan	Iduronic acid/glucuronic acid	Galactosamine	X-sulfated
Keratan	Galactose	Glucosamine	X-sulfated
Hyaluronic acid	Glucuronic acid	Glucosamine	None

GAG- SUB CLASSES

Category	Modifications	Subclasses
Heparan	X-sulfated Y-acetylated/sulfated	HS-0S, HS-NS , HS-6S, HS-triS, HS-diS1
Chondroitin/dermatan	X-sulfated	C4S, C6S, DS
Keratan	X-sulfated	Di-KS, Mono-KS

CS= chondroitin sulfate

Di-KS= di-sulfated keratan sulfate

DS= dermatan sulfate

HS= heparan sulfate

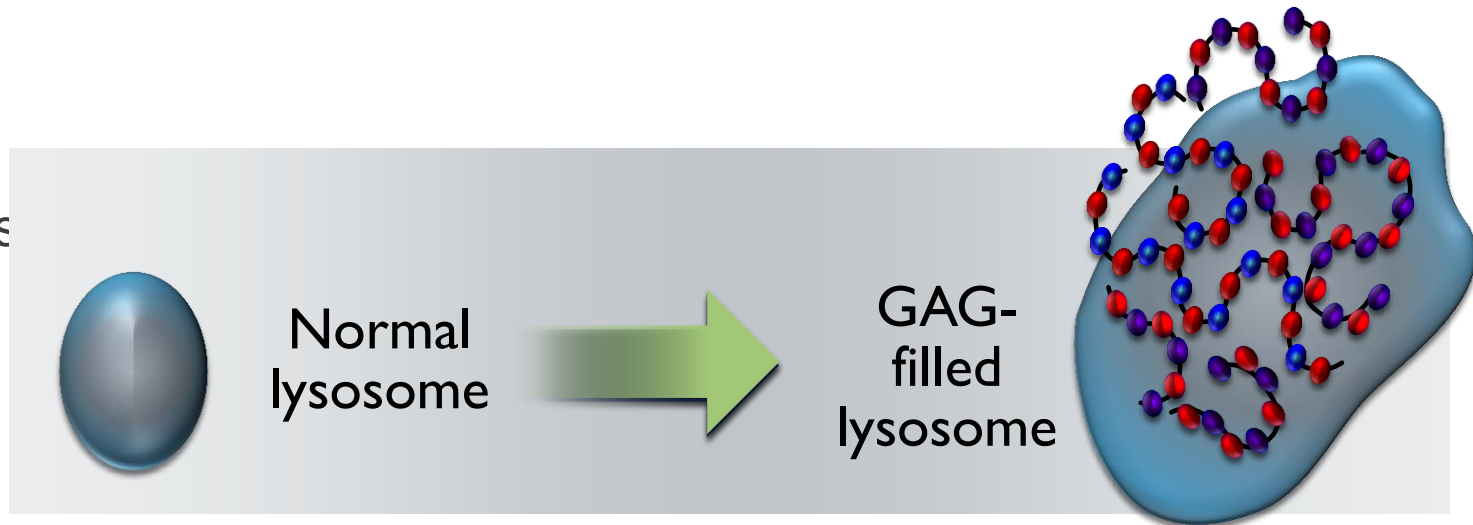
Mono-KS= mono-keratan sulfate

PATHOPHYSIOLOGY

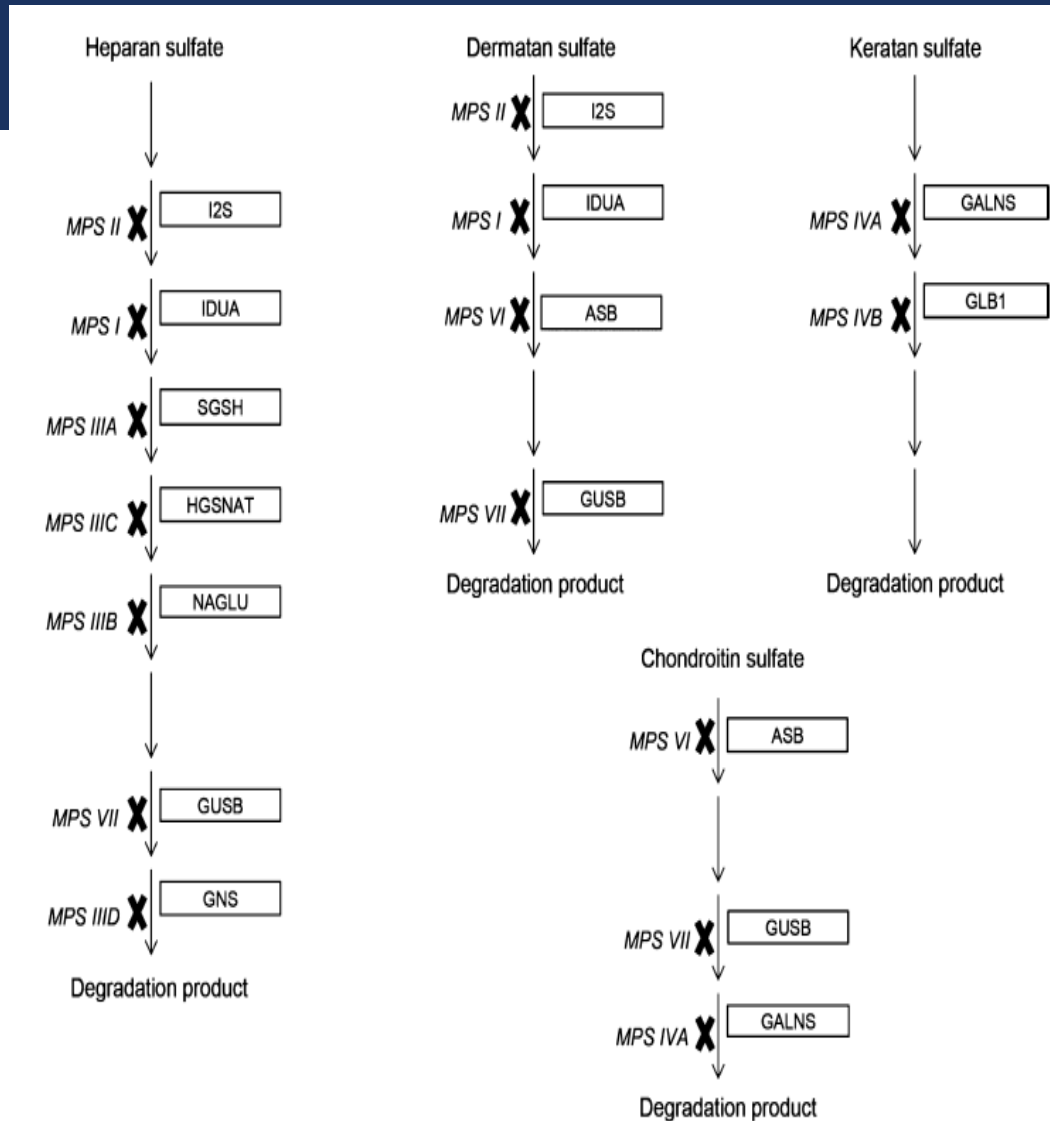
MUCOPOLYSACCHARIDOSES

- Patients with mutations in genes coding for lysosomal enzymes

- Thus



Impaired GAG catabolism leading to MPSs



MPS: BURDEN OF DISEASE

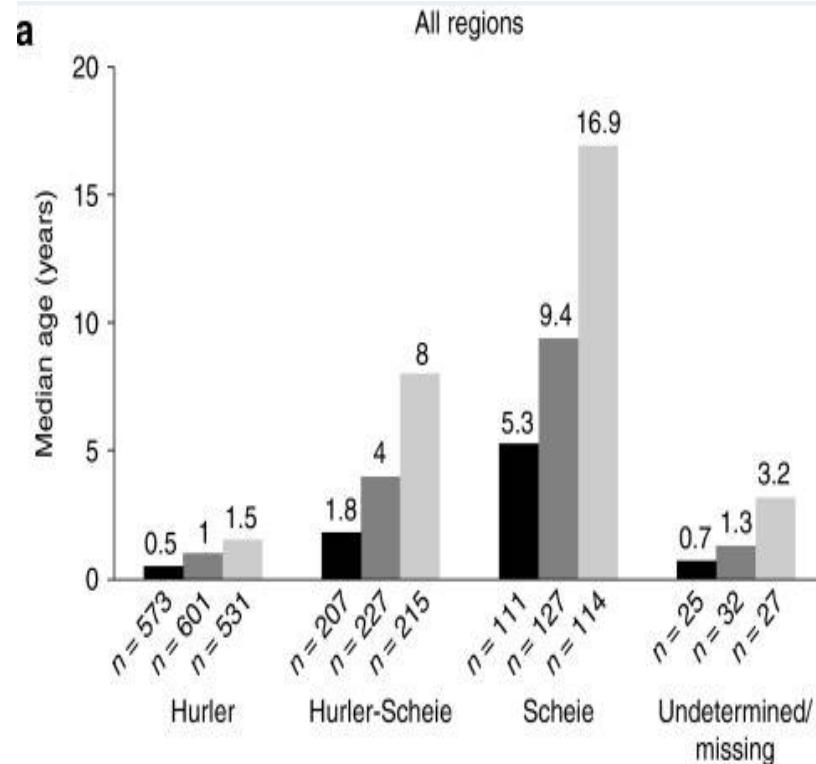
Patients

- GAG accumulation occurs systemically and can result in widespread tissue damage¹
 - Tissue damage is progressive and may be irreversible in some areas of the body^{1,2}
 - Widespread pain, decreased mobility, and decreased respiratory function affect the patient's quality of life³
- Most patients with attenuated forms of MPS are able to receive mainstream education¹
 - Patients with severe forms of MPS experience developmental delay and cognitive decline⁴

Caregivers

- Diagnosis can be obtained by pediatricians or primary care physicians, whereas symptom management often requires multispecialty care⁴
- As mobility decreases, patients may become more dependent on caregivers to assist them with daily activities¹
- Home adaptations may be required for an affected child¹

SYMPTOM ONSET AND DIAGNOSIS TIMELINE

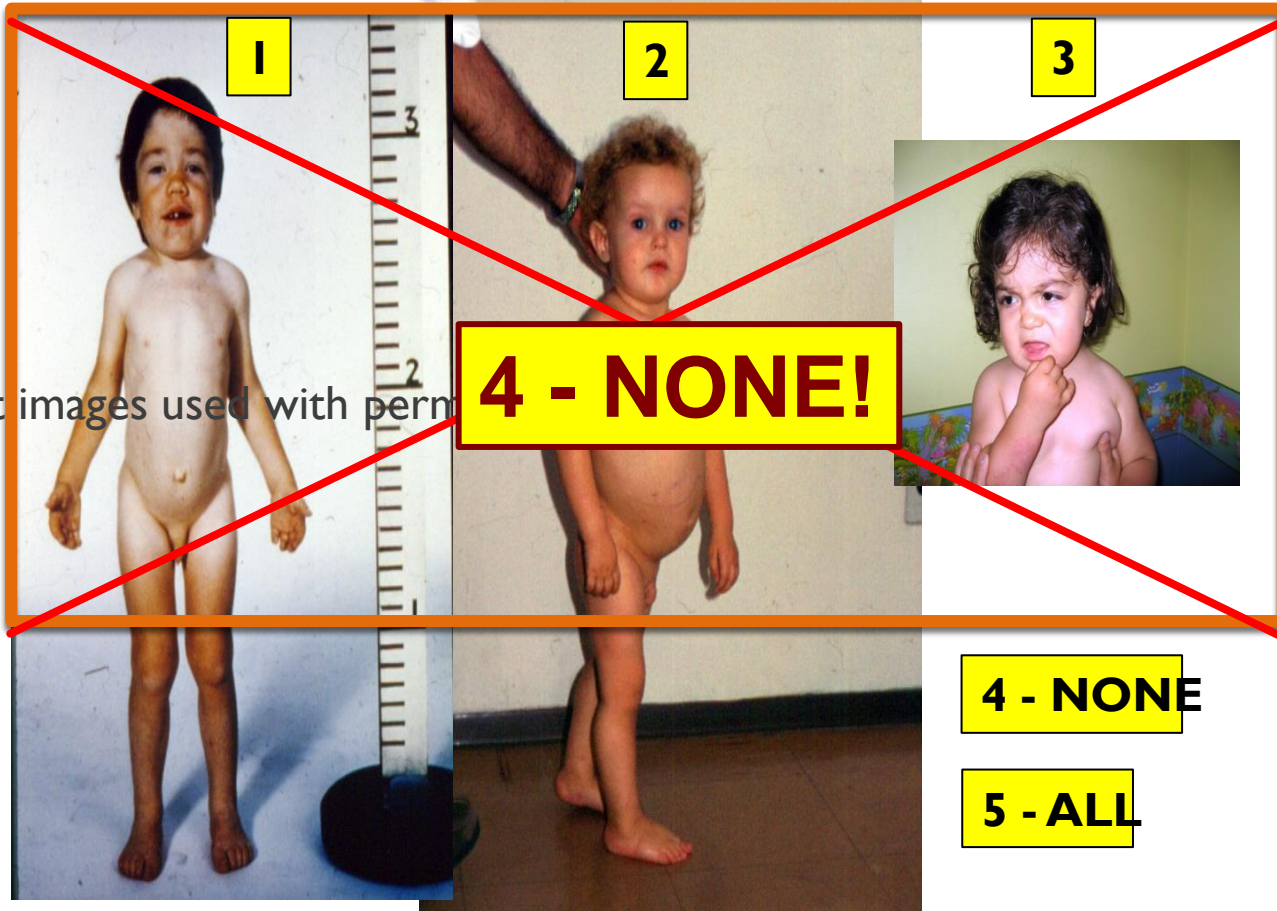


The delay in diagnosis may result in delayed disease management

WHEN SHOULD YOU CONSIDER MPS?

- Child born 'normal', slowly progressive course
- Coarse facies (not always)
- Corneal clouding (not always)
- Hepato- and/or splenomegaly (not always)
- Joint stiffness (not always)
- Bone dysplasia (may be mild)
- (Respiratory infections and/or distress)
- (Heart valve disease)
- (Previous surgery – mainly for hernia, tonsils)
- Cognitive decline (in some cases, may be prominent)

QUIZ 1: WHICH OF THESE PATIENTS HAS MPS?



■ Patient images used with permission

4 - NONE

5 - ALL

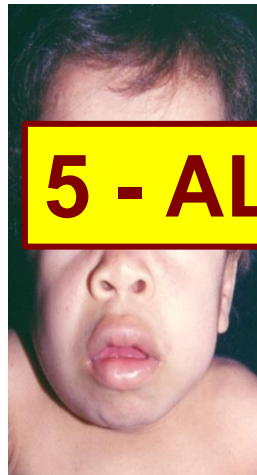
QUIZ 2: WHICH OF THESE PATIENTS HAS MPS?

Patient images used with permission

1



2



5 - ALL!

3



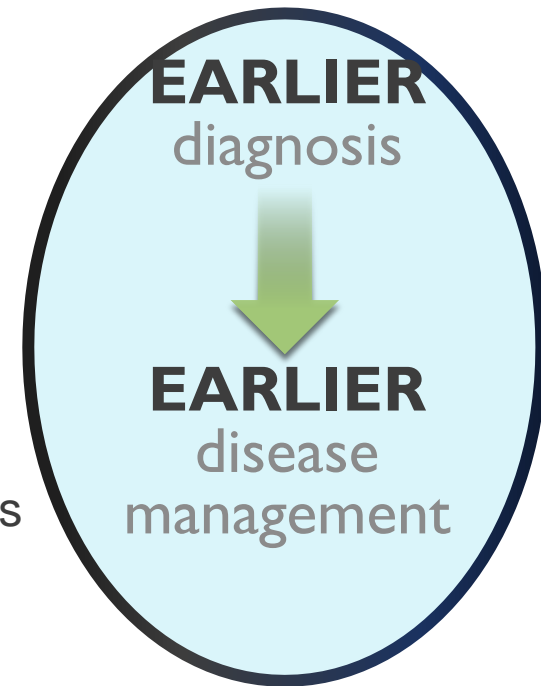
4 - NONE 5 - ALL

1ST-TAKE HOME MESSAGE

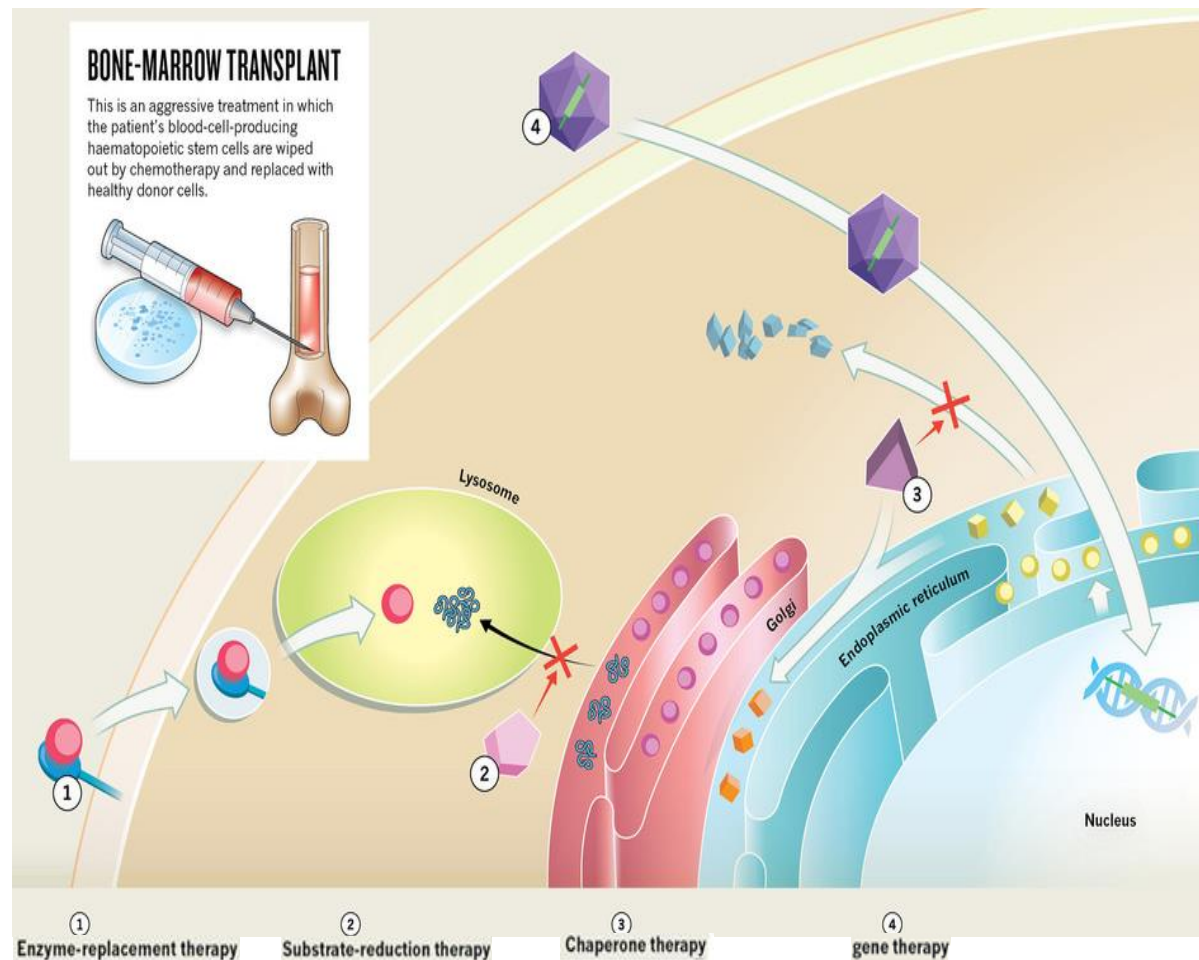
- Clinical suspicion is important, but you need the lab to make a diagnosis of MPS

EARLY DIAGNOSIS IS KEY TO MANAGEMENT OF MPS

- Diagnosis may be based on a unique combination of symptoms rather than on a single presenting symptom
- Patients often visit multiple physicians before receiving a diagnosis of MPS
 - Diagnostic delays are common, particularly in patients with attenuated forms of MPS
- Screening of newborns in high-risk populations is not widespread, but may result in earlier diagnosis



TREATMENTS



TREATMENTS



TREATMENTS



TREATMENTS

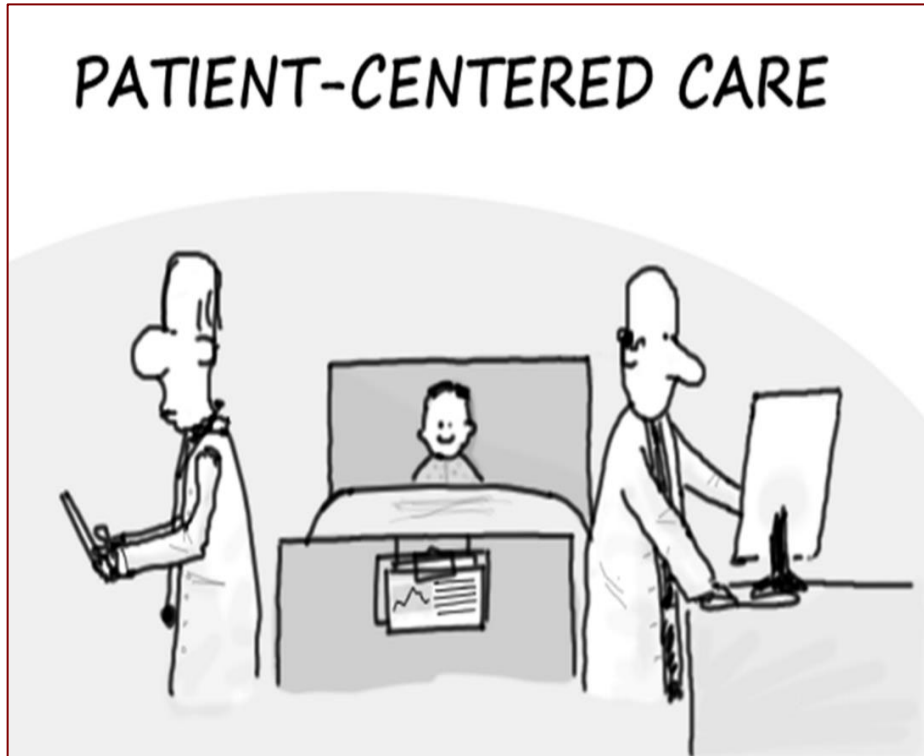


TREATMENT- WHAT IS THE BEST?

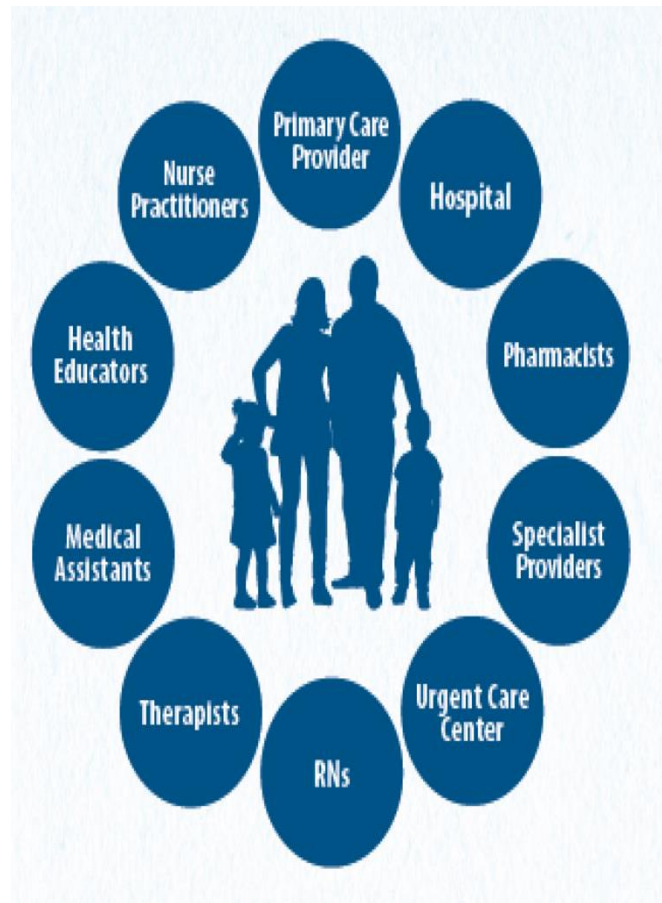
- ✓ Type of MPS
- ✓ Severity of the Phenotype
- ✓ Degree of System/Organ Involvement
 - (CNS, Bone, etc)
- ✓ Age of the Patient

TREATMENT

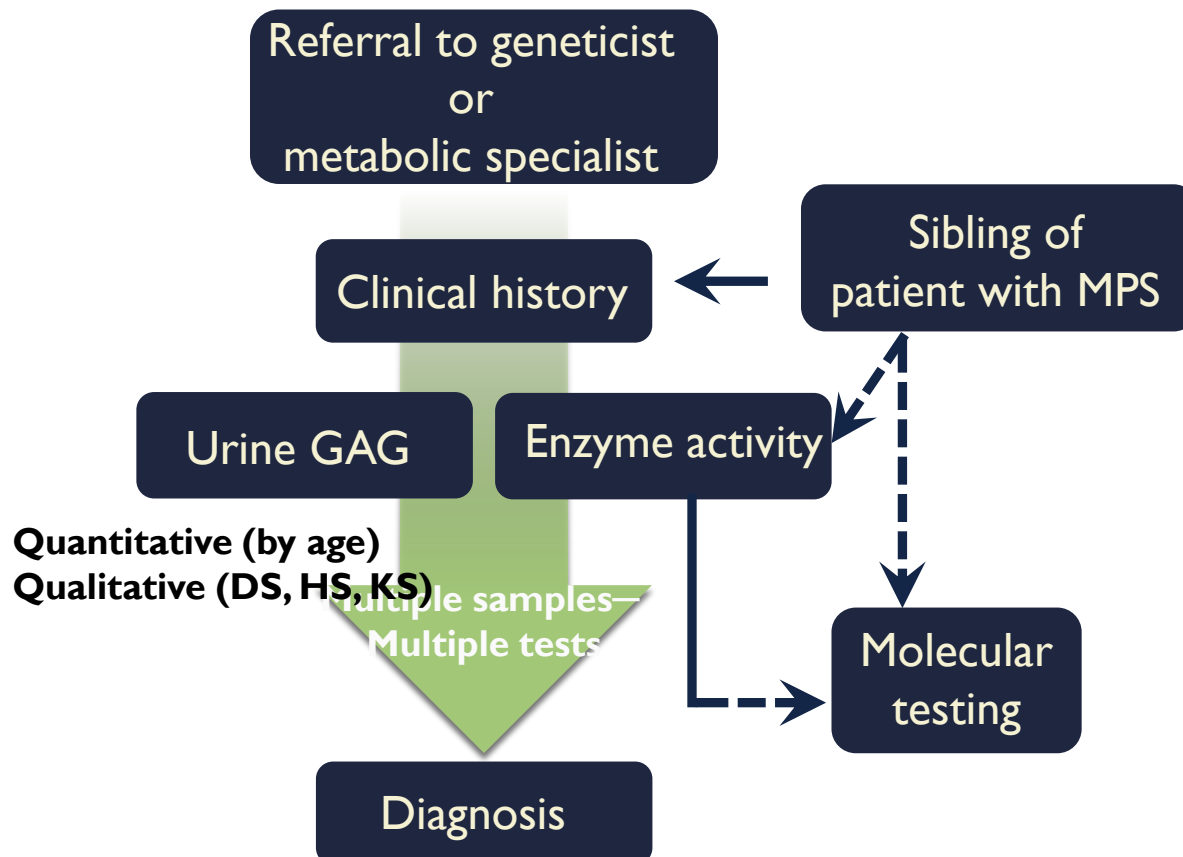
PATIENT-CENTERED CARE



MULTIDISCIPLINARY TEAM



DIAGNOSTIC ALGORITHM



CURRENTLY AVAILABLE DIAGNOSTIC TESTS



- Urinary GAG test
 - Accumulation of GAGs in lysosomes results in elevated GAGs in the urine
 - May be qualitative or quantitative



- Enzyme assays
 - Measures activity of lysosomal enzymes
 - Required for definitive diagnosis
 - Commonly performed on dried blood spots



- Genetic testing
 - Identifies mutation(s)

CLUSTERS IN LATIN AMERICA

- There is high allelic heterogeneity in all MPS genes, with high numbers of genetic mutations reported so far.
- MPS are considered ultra-rare disorders, but clusters of patients can be identified in some areas, usually associated to relative isolation, founder effect and endogamy.
- The aim of this work was to report clusters of MPS in Brazil, Dominican Republic, Ecuador, Haiti and Panama involving the types IIIB, IIIC, IVA and VI.

SAMPLES AND METHODS

- Blood samples were obtained from 28 Ecuadorian patients, 43 Brazilian patients and 7 patients from Dominican Republic.

Diagnoses

- All patients were identified by low enzyme activity and confirmed by molecular analyses (Sanger sequencing [Applied 3500xL] and/or Next-generation sequencing [IonTorrent]).

Bioinformatics analyses

- To predict the possible pathogenicity of the novel variants detected, the Mutation Taster, SIFT and PredictSNP software were used.

RESULTS

Country	State	Number of patients	Type of MPS
Brazil	Paraíba (PB)	7	IIIC
Brazil	Paraíba (PB)	23	IVA
Brazil	Bahia (BA)	13	VI
Dominican Republic	Distrito Nacional	7	VI
Ecuador	Manabi	20	IIIB
Ecuador	Pastaza	8	IVA

RESULTS

Six clusters were identified in Brazil and Ecuador

Figure 1: Map of clusters in Ecuadorian (A), Brazil (B) and Dominican Republic (C)

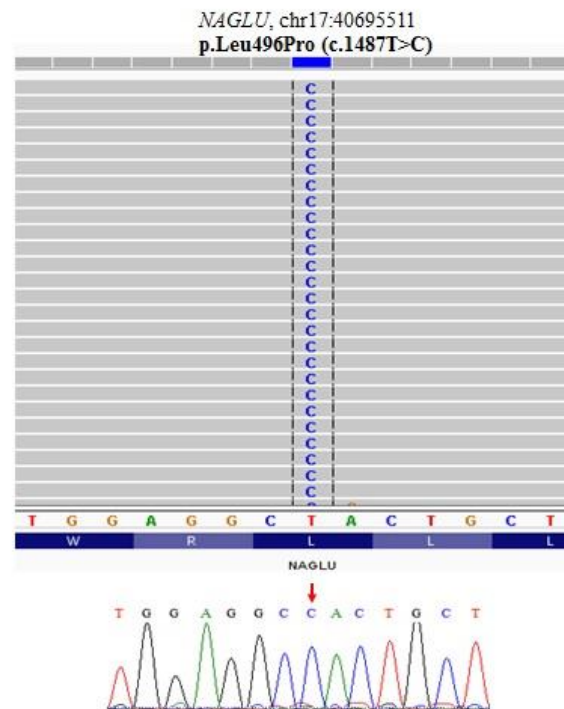


RESULTS

Molecular analyses of MPS IIIB patients from Ecuador

- Molecular analyses identified 17 homozygotes (85%) for the variant p.Leu496Pro, two compound heterozygotes (p.Leu496Pro/Arg482Gly; p.Leu496Pro/?) (10%). Confirmation of the variant by Sanger sequencing (Figure 2).

Figure 2: Next Generation sequencing of *NAGLU* identifying p.Leu496Pro



sequencing of *NAGLU* identifying

RESULTS

Molecular analyses of MPS IIIC from Brazil

- Molecular analyses identified the following variants: c.234+1G>A, c.372-2A>G, c.525dupT, p.Ser541Leu, c.1348delG, p.Asn258Ile (Martins et al. 2014).

Molecular analyses of MPS IVA from Brazil

- Molecular analyses identified 12 homozygotes (52%) and 3 heterozygotes for p.Ser341Arg (13%). All homozygote patients shared a common haplotype within the cluster. Haplotype analyses suggest a common ancestor (Bochernitsan et al., 2018).

Molecular analyses of MPS IVA from Ecuador

- Molecular analyses identified 8 homozygotes (100%) for p.Arg251Ter. This variant has been reported as disease associated (Morrone et al., 2014).

Molecular analyses of MPS VI patients from Brazil

- Molecular analyses identified 13 homozygotes (100%) for the variant p.His178Leu sharing the same haplotype within the cluster. Haplotype analyses suggest a common ancestor (Costa-Motta et al., 2014).

RESULTS

Molecular analyses of MPS VI patients from Dominican Republic

- Molecular analyses identified 7 homozygotes (100%) for the variant p.His178Leu sharing the same haplotype within the cluster. Haplotype analyses suggest a common ancestor (Costa-Motta et al., 2014).

Pathogenicity prediction

- The variant identified in the MPS IIIB patients from Ecuador is classified with uncertain clinical significance by ClinVar. Polyphen prediction suggests that this alteration is probably damaging with conservation of this leucine in several species (Adzhubei et al., 2010).

RESULTS

- All MPS IIIB patients from Ecuador had a very severe phenotype suggesting that p.Leu496Pro in homozygosis could be associated with a severe phenotype (Figure 4).

Figure 4: Two untreated MPS type B patients homozygotes for p.Leu496Pro

a) 9 year old male; b) 10.1 year old male. Red arrow: note hypertrichosis.

Consent obtained for image use.



RESULTS

- Likewise, all MPS IVA patients homozygotes for p.Arg251Ter had a very severe phenotype (Figure 5).

Figure 5: Three untreated siblings with MPS IVA homozygotes for p.Arg251Ter

- a) 4.11 year old male; b) From left to right: 11.4 year old female, 21 year old male and 4.11 year old male. Red arrow: note *pectus carinatum*, genu valgum.

Consent obtained for image use.

A)



B)



DISCUSSION AND CONCLUSIONS

- We report five MPS clusters in Brazil and Ecuador.
- Haplotype analyses were already performed in two clusters, suggesting a common ancestor.
- Our findings indicate that clusters of specific MPS types are not rare in Latin America, and provide an opportunity for targeted health care and community genetics actions, including specific management and general preventive measures.
- These findings are also important for epidemiological studies, genetic counseling and genotype-phenotype correlation.