

Consideraciones genéticas en la enfermedad de Fabry



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Servicio Medicina Interna

DEFINICIÓN GENÉTICA

“La **Genética** es el estudio de la herencia, el proceso en el cual un padre le transmite ciertos genes a sus hijos.”



“La **Genética** es el campo de la biología que busca comprender la herencia biológica que se transmite de generación en generación.

El principal objeto de estudio la Genética son los **genes**, formados por segmentos de ADN (doble cadena) y ARN (cadena simple).”

WIKIPEDIA
The Free Encyclopedia



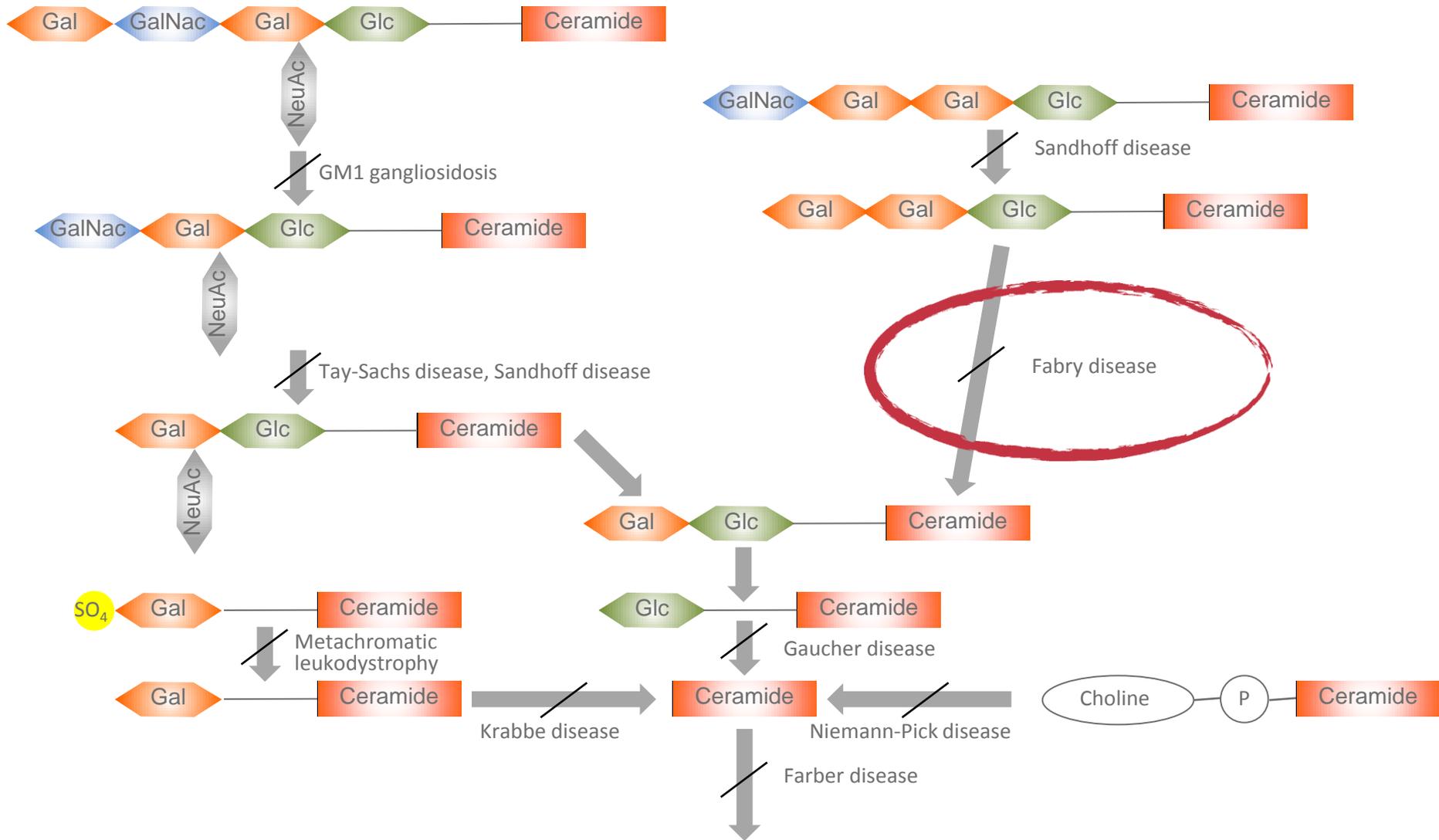
DEFINICIÓN

“La enfermedad de Fabry, también conocida como deficiencia de la alfa-galactosidasa-A, causa una acumulación de material graso en el sistema nervioso autónomo, los ojos, los riñones, y el sistema cardiovascular. La enfermedad de Fabry es la única enfermedad por almacenamiento de lípidos **ligado a X.**”



“La enfermedad de Fabry es una enfermedad de almacenamiento lisosómico **hereditaria ligada al cromosoma X** derivada de **mutaciones en el gen que codifica la enzima α -galactosidasa.**”







Globotriaosylceramide



α -Galactosidase A



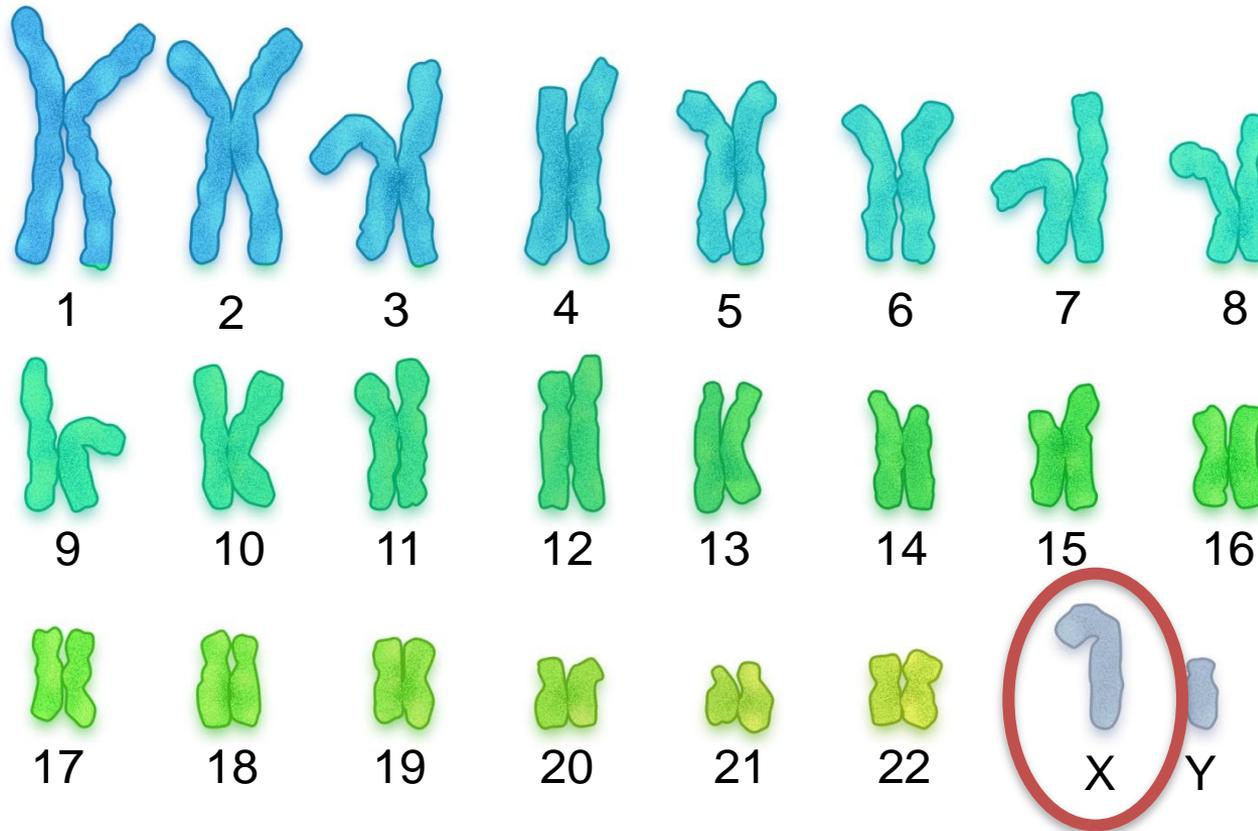
Galactose



Lactosylceramide

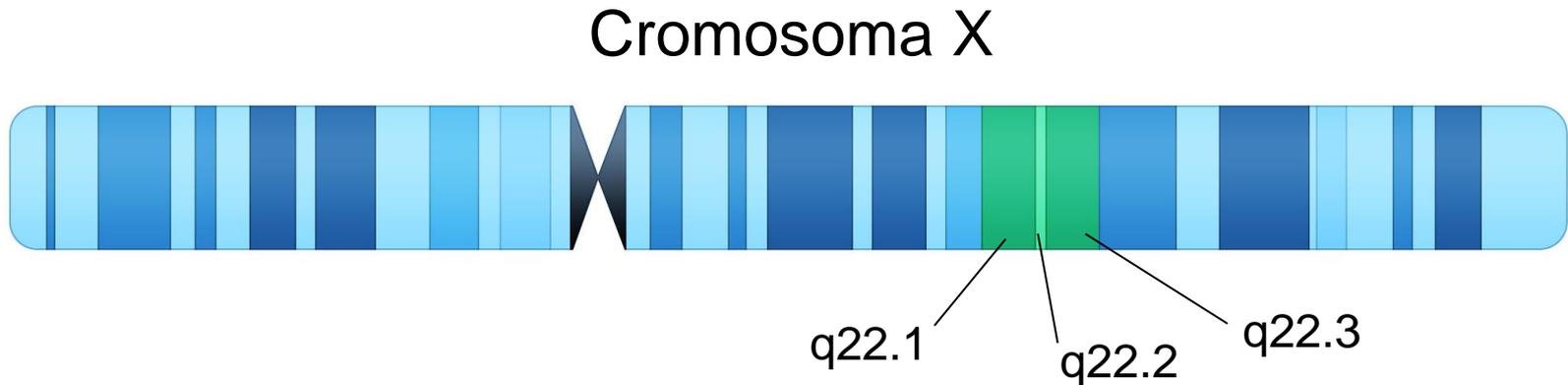
Gen de la alfa-galactosidasa (GLA)

El gen *GLA* se encuentra en el cromosoma X¹



El gen *GLA* en Xq22

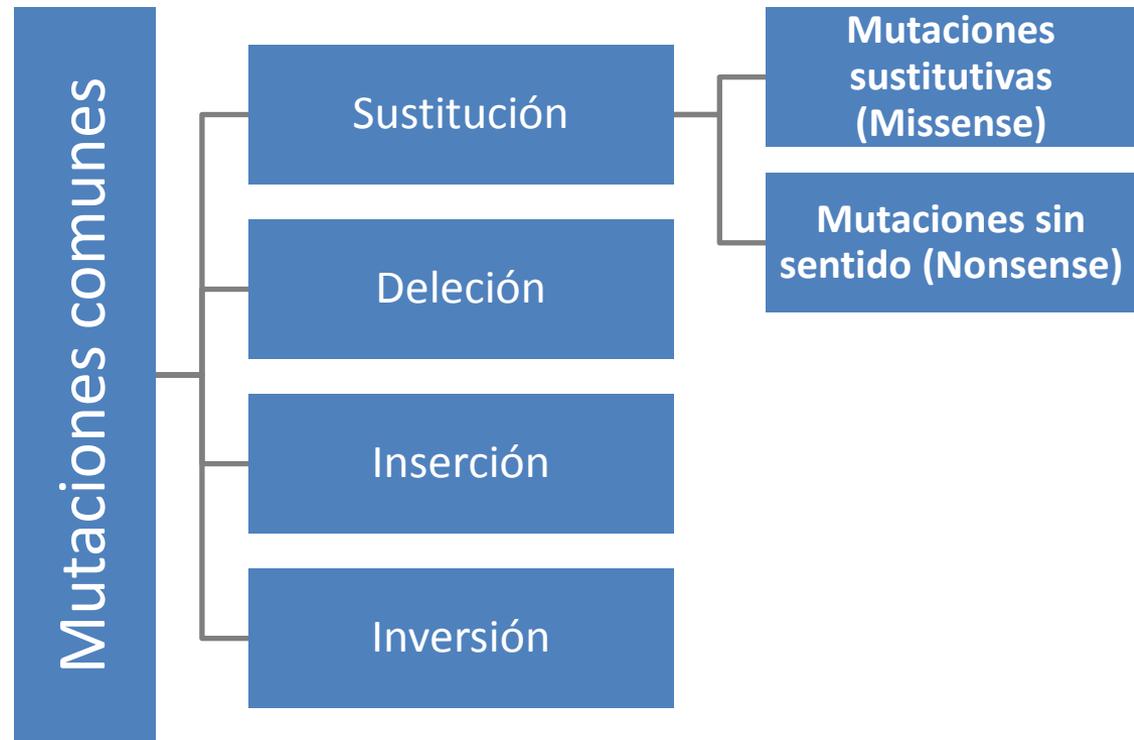
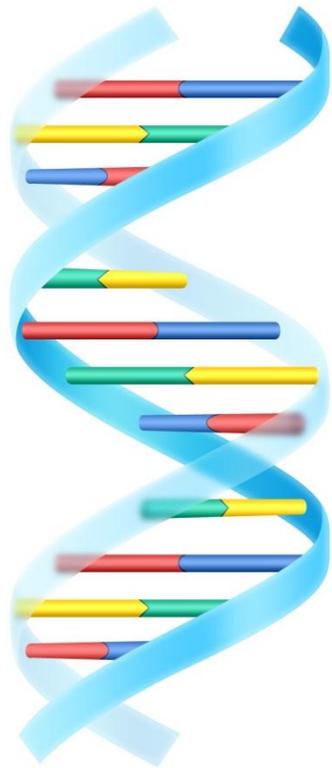
El gen *GLA* se encuentra en la posición 22 del cromosoma X¹



En la enfermedad de Fabry, el gen *GLA* está ausente o es deficiente²

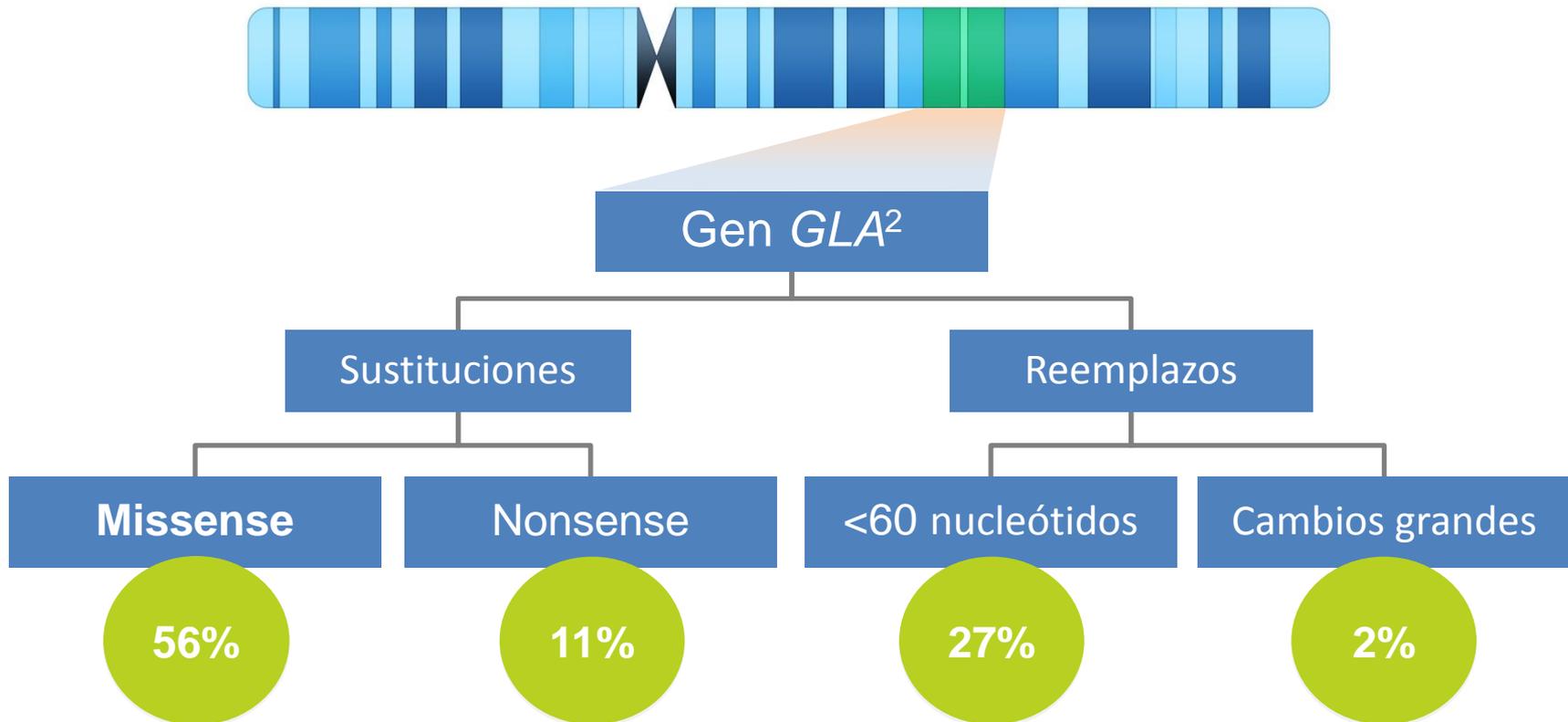
Mutaciones genéticas

Son cambios permanentes en el ADN



Mutaciones genéticas en la enfermedad de Fabry

>500 mutaciones en el gen *GLA* en la enfermedad de Fabry¹



¹Benjamin 2009; ²Gal 2006

MUTACIONES GEN GLA

Biochemical and Molecular Genetic Basis of Fabry Disease

GREGORY M. PASTORES,* YEONG-HAU H. LIEN[†]

**Neurogenetics Program, Department of Neurology and Pediatrics, New York University School of Medicine, New York, New York; and [†]Department of Medicine, University of Arizona Health Sciences Center, Tucson, Arizona.*

J Am Soc Nephrol 13: S130–S133, 2002



| | | | | | | |
|---------|---------|-------|---------|---------|----------|-------|
| M1T | S65T | G128E | C202Y | N215S | Q279E | A377D |
| M1I | E66Q | Y134S | P205T | Y216D | Q279H | W340R |
| P6X | IVS G-T | G138R | W209X | R220X | Q280H | W340X |
| A20P | M72V | C142Y | Ex4-SD | G258R | M284T | R342Q |
| A31V | Y86C | A143T | 613D9 | P259R | W287C | R342X |
| L32P | L89P | S148R | IVS4 +1 | G260A | L294X | R356W |
| P40S | I91T | A156V | IVS4 +2 | N263S | M296I | E358K |
| W44X | D92Y | G163V | IVS4 +4 | V269A | M298K | G361R |
| H46R | C94S | L166V | | N272K | N298H | R363H |
| H46Y | C94Y | D170V | | W277X | N298S | Y365X |
| W47G | W95S | 409D1 | | 239D2 | N301G | G373S |
| R49P | A97V | 520DT | | 240ins2 | R301Q | A377D |
| C52R | R100T | | | 241ins1 | R301X | P409A |
| 125D13 | R112C | | | 256D1 | D313Y | P409T |
| 154DT | F113L | | | 257D2 | N329K | 777D1 |
| DE358 | F113S | | | M267I | G328R | |
| Complex | 304D1 | | | G358R | 898*ins1 | |
| | Complex | | | P259R | 987D1 | |

MUTACIONES GEN GLA

Mutaciones en los 7 exones

93% de los genotipos se corresponden con variante clásica de la EF

18 genotipos asociados con variante cardiaca de la EF.

Gran heterogeneidad en los genotipos

Variante clásica: R220X, R227Q, E398X...

Variante renal: S78X, C126-127, CATG, A352D,...

Variante cardiaca: R112H, R301Q, G328R, R404,...

Lysosomal Disease Network's 10th Annual WORLD Symposium 2014

Monday, February 10, 2014 - Friday, February 14, 2014

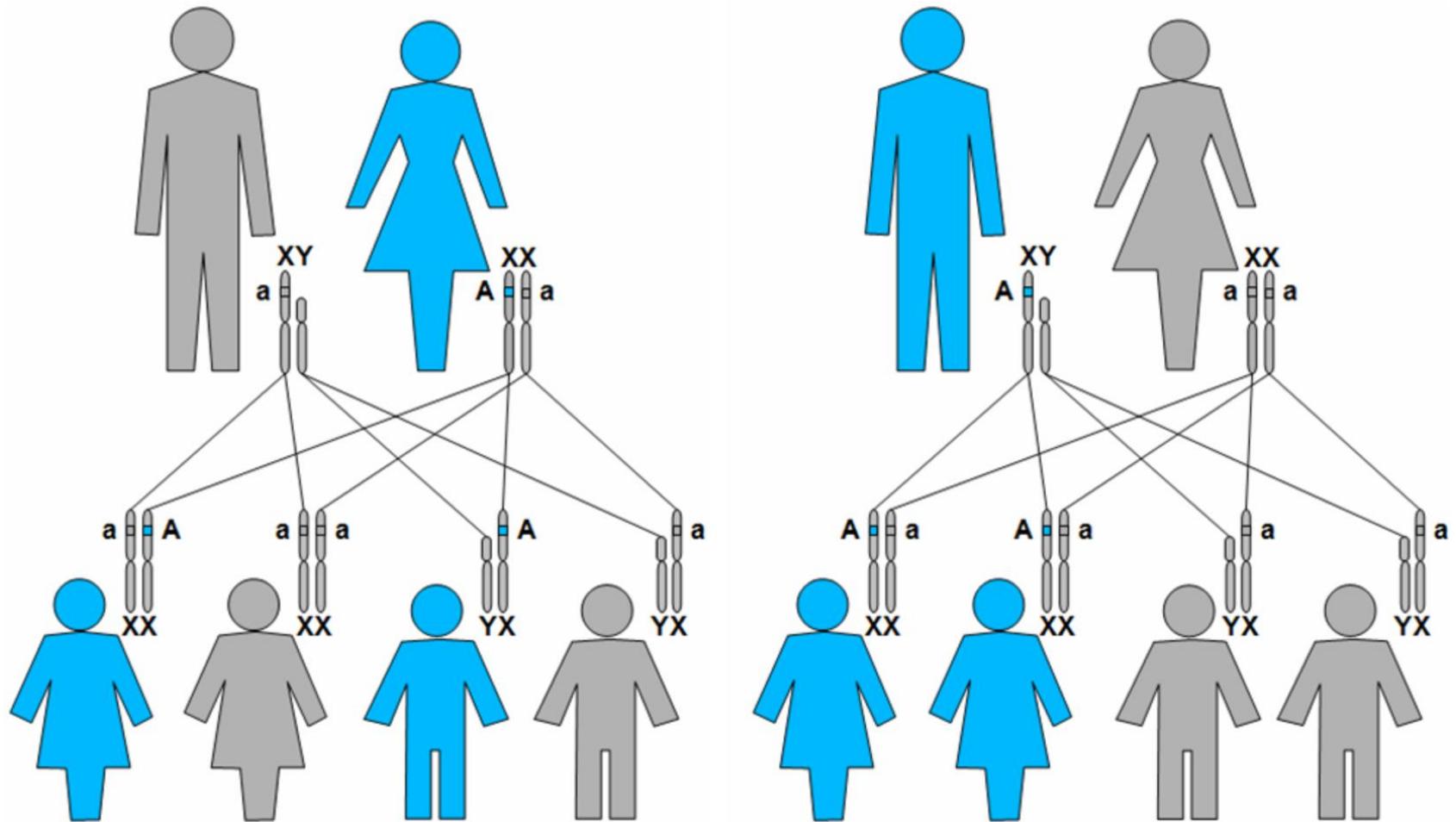
Manchester Grand Hyatt San Diego, San Diego, CA

The demographics and prevalence of the most frequently reported α -galactosidase A mutations in Fabry patients: data from the Fabry Registry

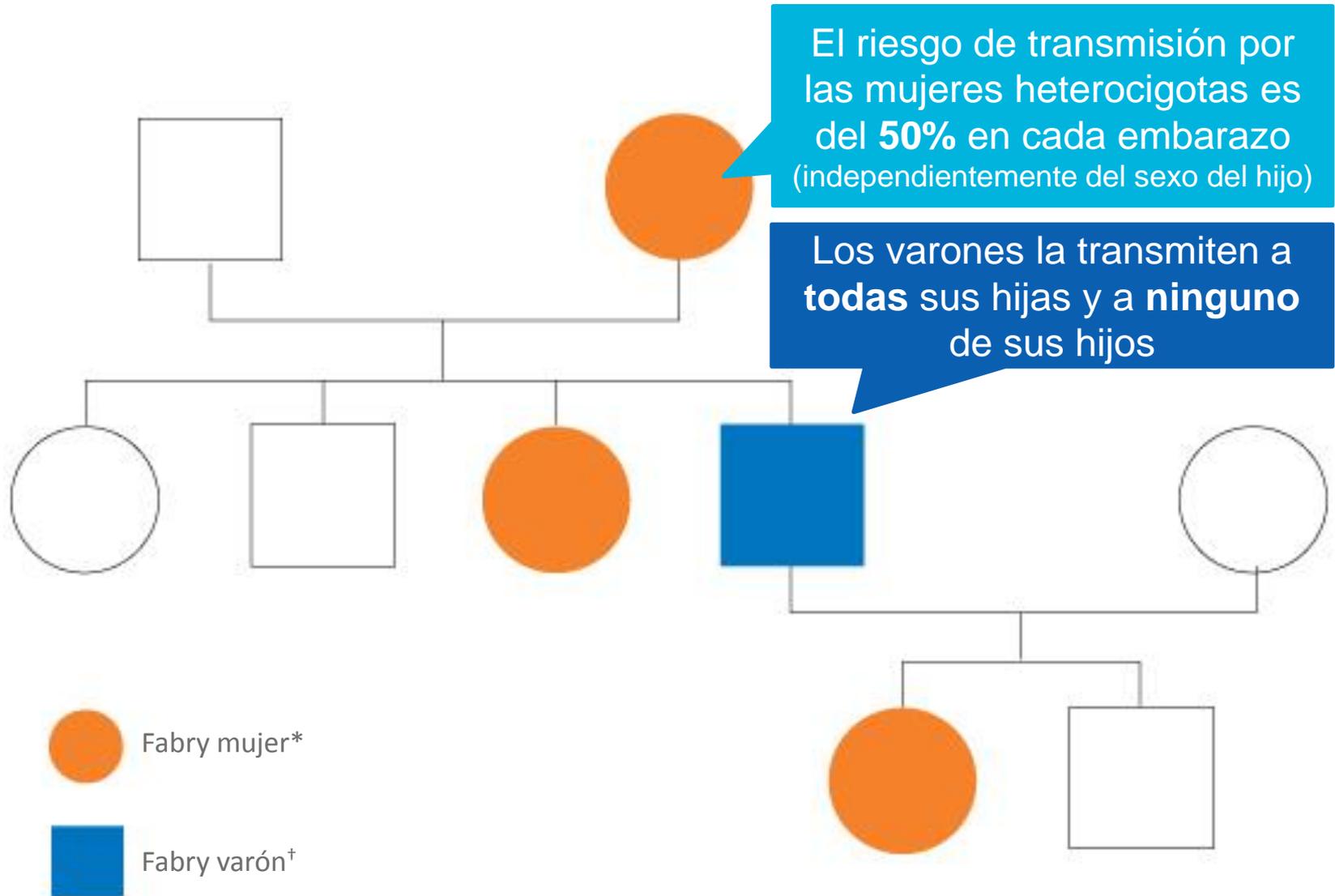
Dawn A. Laney, *Emory, Decatur, GA, USA*

- **Fabry Registry. Genotipo de más de 3200 pacientes**
- **15 mutaciones más repetidas: 10 *missense*.**
- **La más repetida (4,8%): N215S.**
- **La mayoría: mutaciones familiares exclusivas o *de novo*.**
- **Gran heterogeneidad en los genotipos**

Herencia ligada al X (NO RECESIVA)



Herencia ligada al X (NO RECESIVA)

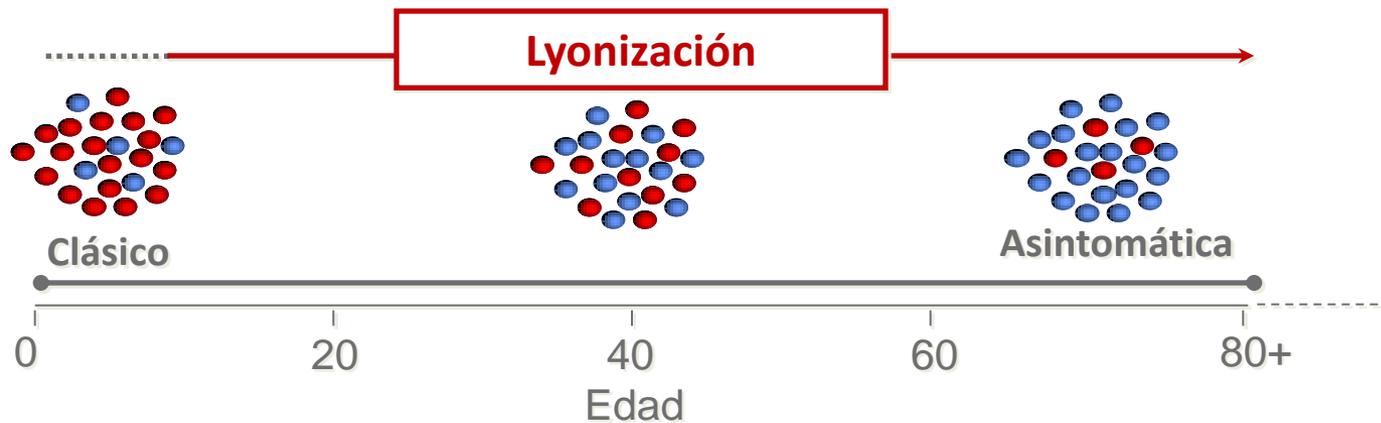


Herencia ligada al X (NO RECESIVA)

Varones hemicigotos (IS: 84%)*

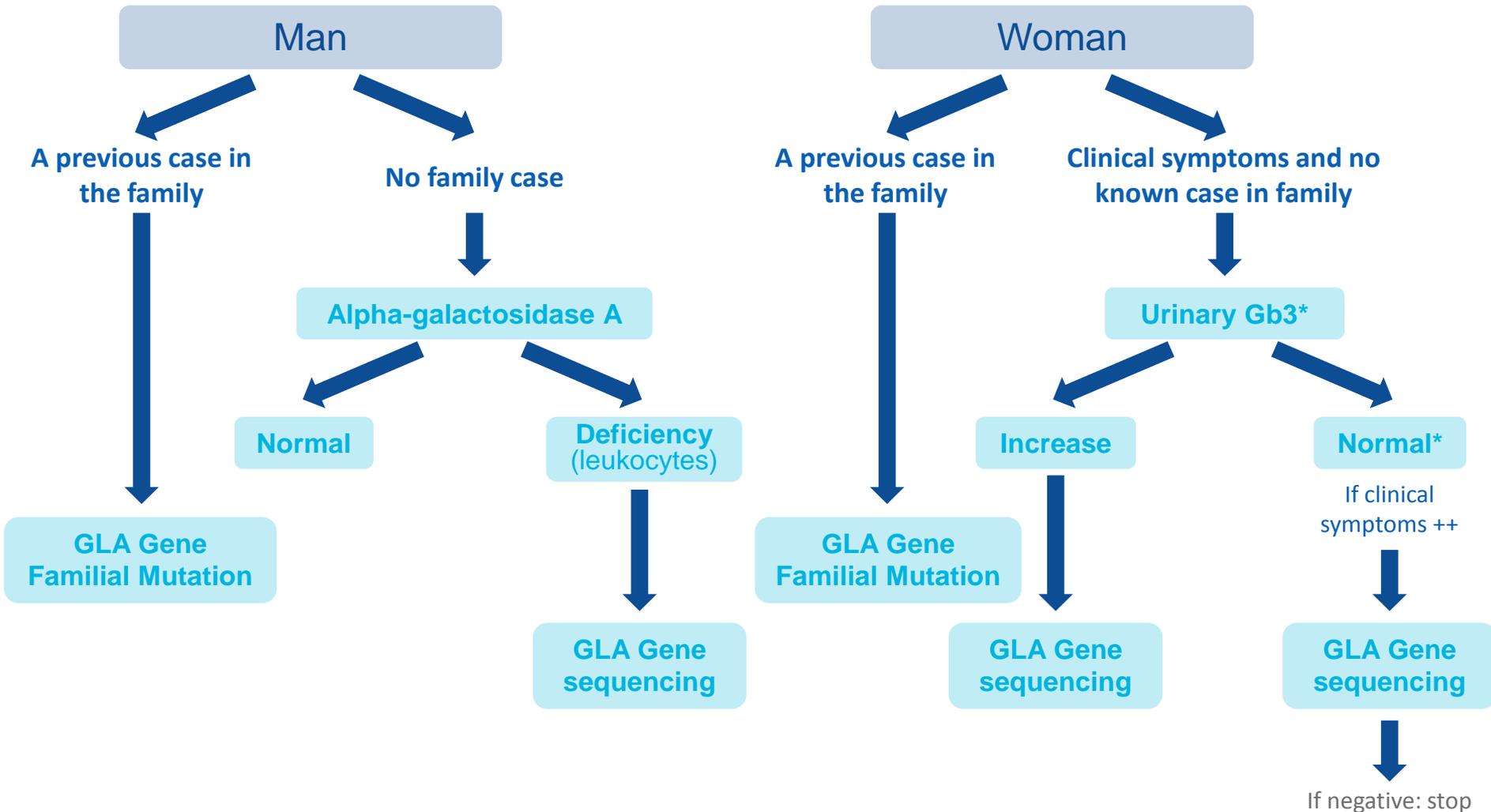


Mujeres heterocigotas (IS: 4%)*



*IS: Índice de Severidad.
Am J Med Genet 2004;129A:136.

¿ESTUDIO ENZIMÁTICO, MOLECULAR,...?



*Urinary Gb3 excretion (men and women) always normal in Fabry disease with certain mutations (and after kidney transplantation).

¿LysoGb3?

How well does urinary lyso-Gb₃ function as a biomarker in Fabry disease?

Christiane Auray-Blais^{a,*}, Aimé Ntwari^a, Joe T.R. Clarke^a, David G. Warnock^b, João Paulo Oliveira^c, Sarah P. Young^d, David S. Millington^d, Daniel G. Bichet^e, Sandra Sirrs^f, Michael L. West^g, Robin Casey^h, Wuh-Liang Hwuⁱ, Joan M. Keutzer^j, X. Kate Zhang^j, René Gagnon^a

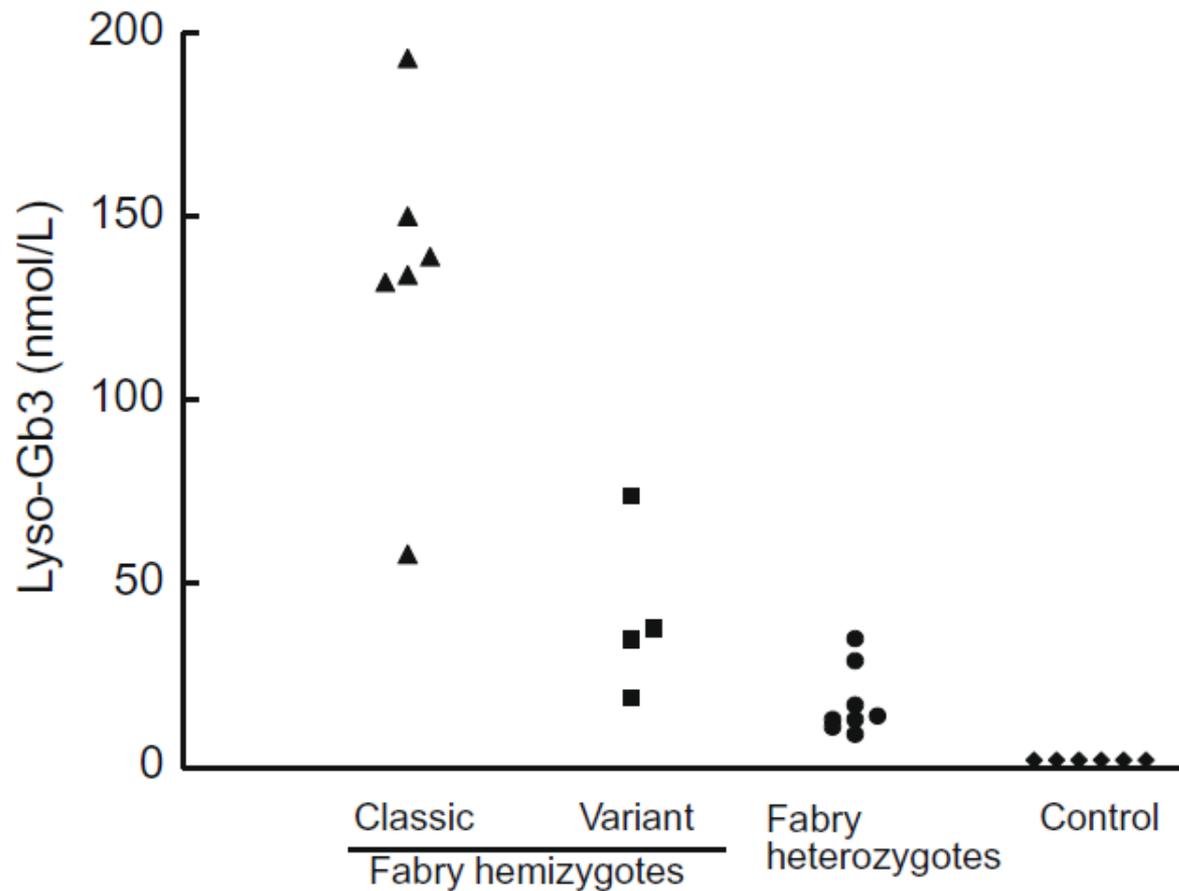
Clinica Chimica Acta 411 (2010) 1906–1914

Plasma globotriaosylsphingosine as a biomarker of Fabry disease

Tadayasu Togawa^a, Takashi Kodama^a, Toshihiro Suzuki^a, Kanako Sugawara^b, Takahiro Tsukimura^a, Toya Ohashi^c, Nobuyuki Ishige^d, Ken Suzuki^d, Teruo Kitagawa^d, Hitoshi Sakuraba^{a,b,*}

Molecular Genetics and Metabolism 100 (2010) 257–261

LysoGb3 como biomarcador



LysoGb3 como biomarcador

- ↑↑↑ en varones forma clásica
- ↑ (respecto controles) en variantes atípicas y mujeres heterocigotas
- Respuesta más sensible que Gb3 al TSE.

Molecular Genetics and Metabolism 100 (2010) 257–261

- Medición, con espectrometría de masas, de Lyso-Gb3 en orina
- Presencia de Lyso-Gb3 en orina es exclusiva de enfermedad de Fabry
- No Lyso-Gb3 en orina de pacientes con diagnóstico molecular de EF (R363H y R118C)

Clinica Chimica Acta 411 (2010) 1906–1914

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The Fabry genotype-phenotype database: an international expert consortium

Robert J. Desnick^a, Dana O. Doheny^b, Dominique P. Germain^c, Yoshikatsu Eto^{d,e}, Frank Weidemann^f, Hitoshi Sakuraba^g, Rong Chen^a, David F. Bishop^a, *^aIcahn School of Medicine at Mount Sinai, New York, NY, USA, ^bIcahn School of Medicine at Mount Sinai, New York, NJ, USA, ^cUniversity of Versailles and Hopital Raymond Poincare, Garches, France, ^dAdvance Clinical Research Center and Asian LSD Center, Kawasaki, Japan, ^eTokyo Jikei University School of Medicine, Tokyo, Japan, ^fUniversity of Wurzburg, Wurzburg, Germany, ^gMeiji Pharmaceutical University, Tokyo, Japan*

- www.hgmd.org, www.fabry-database.org, otros “registros Fabry”
- **School of Medicine at Mount Sinai**
- **Acceso libre.**
- **Implicaciones en seguimiento, relación con el fenotipo, decisión de TSE...**

Original Article

Gene Mutations Versus Clinically Relevant Phenotypes Lyso-Gb3 Defines Fabry Disease

Markus Niemann, MD; Arndt Rolfs, MD; Stefan Störk, MD, PhD; Bart Bijmens, PhD;
Frank Breunig, MD; Meinrad Beer, MD; Georg Ertl, MD; Christoph Wanner, MD;
Frank Weidemann, MD

Background—Currently, no method is available to identify α -galactosidase A (agalA) mutations determining clinically relevant Fabry disease. In our largest European Fabry cohort, we investigated whether a biomarker, specific for the defect, could stratify persons at risk.

Methods and Results—A total of 124 individuals with agalA mutations were investigated with a comprehensive clinical workup, genetic analysis, and laboratory testing, including measurements of agalA activity and lyso-Gb3 (degradation product of the accumulating Gb3). Additionally, an extensive family screening with a clinical workup of relatives was performed. The patient population was divided into 2 samples: previously described mutations (n=72) and novel mutations (n=52). The patients with previously described mutations were subdivided into 2 groups: classical mutations, which were known to cause the classic type of Fabry disease with specific symptoms and a high risk for major events in all 3 main organs (heart, kidney, and central nervous system), and atypical mutations without the typical presentation. All patients with atypical mutations (n=17) had lower lyso-Gb3 levels than any of the patients with classical Fabry disease (n=55). A cutoff value of 2.7 ng/mL separated the 2 groups. Six out of 52 patients with novel mutations showed a lyso-Gb3 level <2.7 ng/mL. Clinical investigation, blinded to lyso-Gb3 results, revealed no classic organ involvement in these patients or their relatives. In contrast, the characterization of patients with lyso-Gb3 \geq 2.7 ng/mL suggested classical Fabry mutations in most of the patients (93%).

Conclusions—Our data show that the biomarker lyso-Gb3 may identify the clinically relevant agalA mutations leading to Fabry disease. (*Circ Cardiovasc Genet.* 2014;7:8-16.)

GENOTIPO y LysoGb3

GENOTIPO:

- Mutación clásica:

Afectación de al menos 2 de 3 órganos (corazón, riñón, SNC)

Síntomas específicos cualquier órgano + 1 evento mayor (IRC est III-IV, HVI severa, ictus)

- Variantes atípicas: resto

124 pacientes con genotipo de EF. De ellos, 23 pacientes con Lyso-Gb3 < 2,7 ng/mL:

17/23 mujeres

5/23 con polimorfismos

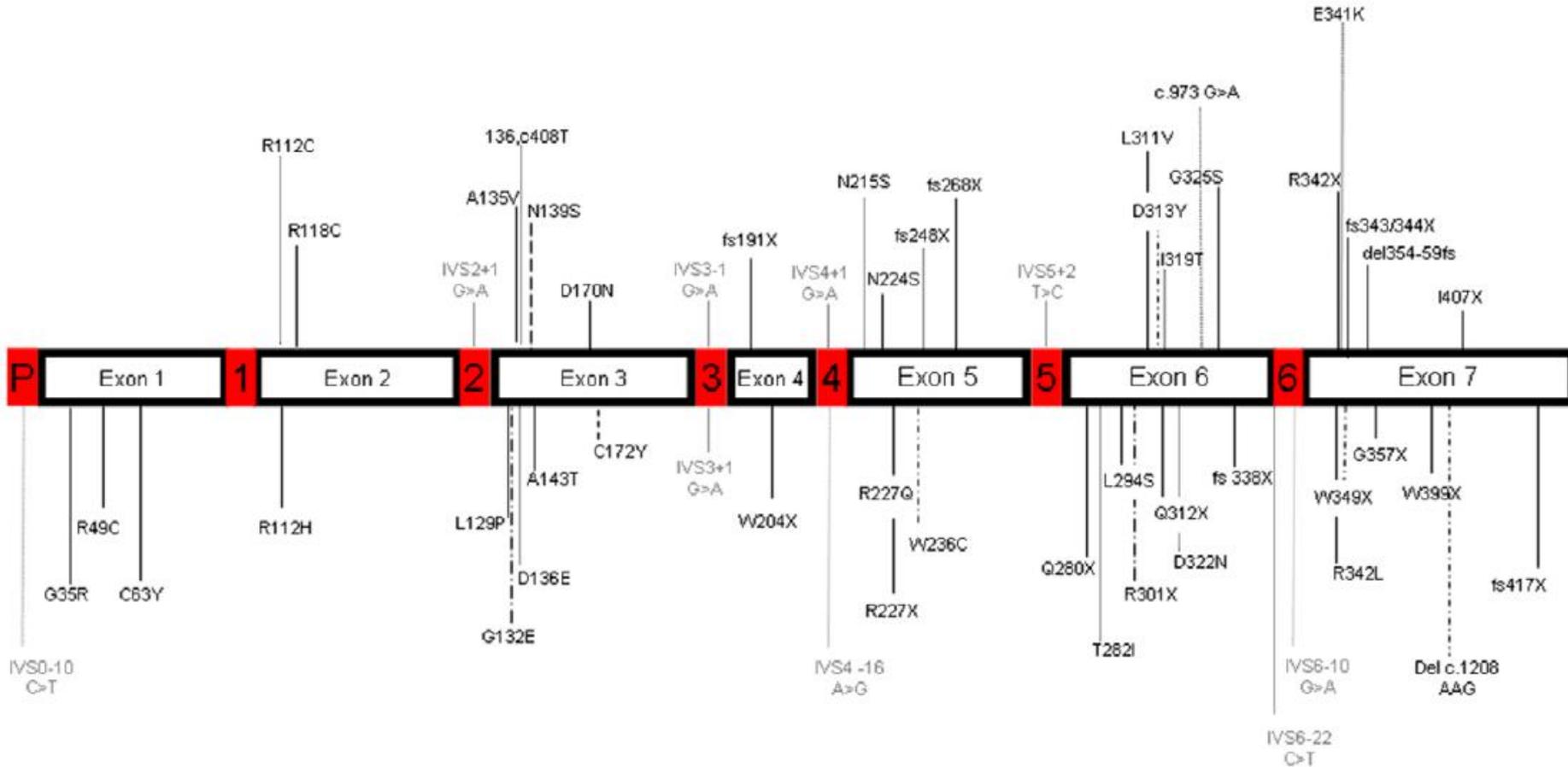
0/23 HVI

2/23 IRC (1 caso HTA, 1 caso sarcoidosis renal)

3/23 ictus (1 cardioembólico, 2 criptogénicos)

Circ Cardiovasc Genet. 2014;7:8-16.

GENOTIPO y LysoGb3



Circ Cardiovasc Genet. 2014;7:8-16.

GENOTIPO y LysoGb3

Hombres vs Mujeres

Table 1. Clinical Baseline Characteristics of the Patients With agalA Mutations Subdivided by Sex

| | Men n=51 | Women n=73 | PValue |
|---|-------------|---------------|---------|
| Age, y | 39±13 | 41±14 | 0.30 |
| Alpha-galactosidase activity, nmol/min per mg protein | 0.06±0.11 | 0.32±0.36 | <0.0001 |
| Lyso-Gb3, ng/mL | 67.8±35.9 | 6.3±4.7 | <0.0001 |

Mutaciones típicas vs Atípicas

Table 3. Clinical Baseline Characteristics of All Patients With Typical and Atypical agalA Mutations

| | Typical n=101 | Atypical n=23 | PValue |
|---|-----------------|----------------|--------|
| Men/women | 45/56 (45%,55%) | 6/17 (24%,76%) | 0.08 |
| Age, y | 40±14 | 40±14 | 0.95 |
| Alpha-galactosidase activity, nmol/min per mg protein | 0.2±0.1 | 0.5±0.6 | 0.02 |
| Lyso-Gb3, ng/mL | 38.7±39.1 | 0.8±0.9 | <0.001 |

Mutaciones típicas vs Atípicas

Table 3. Clinical Baseline Characteristics of All Patients With Typical and Atypical *agaA* Mutations

| | Typical n=101 | Atypical n=23 | P Value | |
|------------------------------|---------------|---------------|---------|---|
| General symptoms | | | | |
| Abnormal sweating (n) | 51 (51%) | 2 (9%) | <0.001 | ● |
| Heat or cold intolerance (n) | 58 (58%) | 4 (17%) | <0.001 | ● |
| Chronic diarrhea (n) | 32 (32%) | 4 (17%) | 0.13 | |
| Sudden deafness (n) | 21 (21%) | 2 (9%) | 0.15 | |
| Angiokeratomata (n) | 41 (41%) | 0 | <0.001 | ● |
| Neurology | | | | |
| Stroke (n) | 11 (11%) | 3 (13%) | 0.50 | |
| Chronic pain syndrome (n) | 33 (33%) | 1 (4%) | 0.003 | ● |
| Acroparaesthesia (n) | 68 (68%) | 4 (17%) | <0.001 | ● |
| Heart | | | | |
| Cardiomyopathy (n) | 49 (49%) | 0 (0%) | <0.001 | ● |
| LV mass, g | 153±59 | 100±28 | <0.001 | ● |
| Kidney | | | | |
| Kidney transplantation (n) | 7 (7%) | 0 (0%) | 0.22 | |
| Dialysis (n) | 6 (6%) | 0 (0%) | 0.28 | |
| Serum creatinin, mg/dL | 1.1±1.3 | 0.8±0.2 | 0.28 | |
| Serum urea, mg/dL | 31±19 | 27±7 | 0.33 | |
| DTPA clearance, ml/min | 91±36 | 111±23 | 0.003 | ● |
| Medication | | | | |
| Beta blocker (n) | 16 (16%) | 4 (17%) | 0.53 | |
| Calcium channel blocker (n) | 3 (3%) | 1 (4%) | 0.57 | |
| ACE inhibitor (n) | 33 (33%) | 5 (22%) | 0.22 | |
| Aspirin (n) | 22 (22%) | 4 (17%) | 0.44 | |
| Disease severity score | 9±6 | 4±4 | <0.001 | ● |

GENOTIPO y LysoGb3

CLINICAL PERSPECTIVE

The current study suggests that the biomarker lyso-Gb3 can differentiate classical Fabry disease from atypical agalA mutations. This is especially helpful when the clinical information about a new mutation is limited and in women patients with an agalA activity level in the gray zone. The data question the current practice to establish the diagnosis of Fabry disease: it seems that the measurement of agalA activity in combination with genotyping is not sufficient for diagnosing Fabry disease. Fabry disease might be redefined based on three criteria: (1) information about the agalA mutation, (2) the level of lyso-Gb3, and (3) the typical Fabry symptoms and organ involvement. If a patient has an agalA mutation and an elevated lyso-Gb3 level and typical symptoms or organ involvement, this combination may be labeled classical Fabry disease. In patients exhibiting an agalA mutation plus elevated lyso-Gb3 levels who are still asymptomatic and free of organ involvement, this constellation may be labeled classical Fabry mutation. Finally, in patients with an agalA mutation but a normal lyso-Gb3 level, it may be labeled atypical α -galactosidase A mutation.

Circ Cardiovasc Genet. 2014;7:8-16.

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A case series study on diagnostic dilemmas in Fabry disease

Bouwien Smid^a, Carla Hollak^a, Ben Poorthuis^a, Janneke Timmermans^b, Gabor Linthorst^a, ^a*Academic Medical Centre, Amsterdam, Netherlands*, ^b*UMC St. Radboud, Nijmegen, Netherlands*

conclude, a misdiagnosis of FD may occur in subjects with a GLA variant and isolated non specific findings, such as HCM or stroke. To prevent misdiagnoses, inappropriate counselling and costly therapy, a structured diagnostic approach is mandatory. Other causes need to be excluded, including concomitant disease and risk factors and the use of medication causing similar inclusion bodies on histology. Certain GLA variants might be interpreted as risk factors for developing FD complications rather than directly disease causing and decisions for treatment should be made with caution. Determination of plasma lysoGb3 might be an important tool to discriminate between non Fabry and non-classical FD patients.

CONSEJO GENÉTICO

J Genet Counsel (2013) 22:555–564

DOI 10.1007/s10897-013-9613-3

PROFESSIONAL ISSUES

Fabry Disease Practice Guidelines: Recommendations of the National Society of Genetic Counselors

Dawn A. Laney • Robin L. Bennett • Virginia Clarke •
Angela Fox • Robert J. Hopkin • Jack Johnson •
Erin O'Rourke • Katherine Sims • Gerald Walter

Mutaciones: <http://www.hgmd.cf.ac.uk/ac/index.php>

- **Aprox 500 mutaciones descritas.**
- **La mayoría, únicas (“mutaciones privadas”)**

CONSEJO GENÉTICO

- 1. Remitir a especialista en metabolopatías y consejo genético**
- 2. Riesgo de afectación familiar**
- 3. Elaboración árbol genealógico completo**
- 4. Identificación familiares “en riesgo”**
- 5. Evaluación preconcepcional (especialista clínico)**
- 6. Opciones:**
 - Donantes de gametos / adopción**
 - Tests genéticos (preimplantacionales, bx coriónica, amniocentesis)**

TAKE-HOME MESSAGES

- Herencia lig X (lyonización)
- Gran heterogeneidad en las mutaciones
- Predominan mutaciones *missense*
- Genotipo y Lyso-Gb3: implicación clínica
- Lyso-Gb3 como biomarcador de enfermedad
- Consejo genético especializado

