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Wladimir Mauhin, Olivier Benveniste, Damien Amelin, Clémence Montagner, Foudil Lamari, et al.. Cornea verticillata and acroparesthesia efficiently discriminate clusters of severity in Fabry disease. PLoS ONE, 2020, 15 (5), pp.e0233460. 10.1371/journal.pone.0233460. hal-02871577

HAL Id: hal-02871577 https://hal.sorbonne-universite.fr/hal-02871577

Submitted on 17 Jun 2020

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Citation: Mauhin W, Benveniste O, Amelin D, Montagner C, Lamari F, Caillaud C, et al. (2020) Cornea verticillata and acroparesthesia efficiently discriminate clusters of severity in Fabry disease. PLoS ONE 15(5): e0233460. https://doi.org/10.1371/journal.pone.0233460

Editor: Maria Vittoria Cubellis, University of Naples Federico II, ITALY

Received: December 31, 2019

Accepted: May 5, 2020 **Published:** May 22, 2020

Peer Review History: PLOS recognizes the benefits of transparency in the peer review process; therefore, we enable the publication of all of the content of peer review and author responses alongside final, published articles. The editorial history of this article is available here: https://doi.org/10.1371/journal.pone.0233460

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Data Availability Statement: All relevant data are within the manuscript and its Supporting Information files.

RESEARCH ARTICLE

Cornea verticillata and acroparesthesia efficiently discriminate clusters of severity in Fabry disease

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Abstract

Backgroud

Fabry disease (OMIM #301 500), the most prevalent lysosomal storage disease, is caused by enzymatic defects in alpha-galactosidase A (*GLA* gene; Xq22.1). Fabry disease has historically been characterized by progressive renal failure, early stroke and hypertrophic cardiomyopathy, with a diminished life expectancy. A nonclassical phenotype has been described with an almost exclusive cardiac involvement. Specific therapies with enzyme substitution or chaperone molecules are now available depending on the mutation carried.

Funding: The author(s) received no specific funding for this work.

Competing interests: WM received honoraria, congress fees and travel assistance from Shire-Takeda, Amicus and Sanofi- Genzyme. OB, DA, CM declare no conflict of interest. FLam has received travel support from Amicus Therapeutics., Shire and Sanofi-Genzyme. He received lecture fees from Actelion Pharmaceuticals. BD has received honoraria from Amicus (member of the scientific board) and Novartis (lectures) and travel fees from Genzyme-Sanofi. VLS has received travel fees and accommodations from Shire and Sanofi-Genzyme. CC has received consultant honoraria and congress fees from Biomarin and Sanofi-Genzyme and has participated in editorial activity with Takeda-Shire. CD has received travel assistance from Shire, Sanofi-Genzyme, Sobi, Orphan Europe, Nutricia, Lucane Pharma, Amicus, and Ultragenyx and honoraria from Amicus and has participated on boards with Ultragenyx and Sanofi. OB, AD, PDH declare no conflict of interest EN has received travel fees from Shire and Sanofi-Genzyme and an honorarium from Amicus. TZ has received congress fees and travel assistance from Sanofi-Genzyme. MM declare no conflict of interest FM has received honoraria from Shire and travel assistance from Sanofi-Genzyme, KHL declare no conflict of interestGB has received travel assistance from Shire, Genzyme-Sanofi and Amicus. GB has received travel assistance from Shire, Genzyme-Sanofi and Amicus. MW and FLab declare no conflict of interest AM has received travel fees and accommodations from Shire, Sanofi-Genzyme and Amicus. CL has received honoraria from Sanofi-Genzyme and travel assistance from Sanofi-Genzyme and Shire. DL has received honoraria and travel assistance from Sanofi-Genzyme and has participated on boards with Amicus. HM received honoraria and travel assistance from Sanofi-Genzyme and Amicus and has participated on boards with Amicus and Shire. OL has received travel support and lecture fees from Amicus Therapeutics, Shire, and Sanofi- Genzyme. This does not alter our adherence to PLOS ONE policies on sharing data and materials.

Numerous clinical and fundamental studies have been conducted without stratifying patients by phenotype or severity, despite different prognoses and possible different pathophysiologies. We aimed to identify a simple and clinically relevant way to classify and stratify patients according to their disease severity.

Methods

Based on data from the *French Fabry Biobank and Registry* (FFABRY; n = 104; 54 males), we applied unsupervised multivariate statistics to determine clusters of patients and identify clinical criteria that would allow an effective classification of adult patients. Thanks to these criteria and empirical clinical considerations we secondly elaborate a new score that allow the severity stratification of patients.

Results

We observed that the absence of acroparesthesia or cornea verticillata is sufficient to classify males as having the nonclassical phenotype. We did not identify criteria that significantly cluster female patients. The classical phenotype was associated with a higher risk of severe renal (HR = 35.1; p < 10^{-3}) and cardiac events (HR = 4.8; p = 0.008) and a trend toward a higher risk of severe neurological events (HR = 7.7; p = 0.08) compared to nonclassical males. Our simple, rapid and clinically-relevant FFABRY score gave concordant results with the validated MSSI.

Conclusion

Acroparesthesia and cornea verticillata are simple clinical criteria that efficiently stratify Fabry patients, defining 3 different groups: females and males with nonclassical and classical phenotypes of significantly different severity. The FFABRY score allows severity stratification of Fabry patients.

Introduction

Fabry disease (FD; OMIM #301 500) is an X-linked lysosomal storage disease caused by an enzymatic defect of the hydrolase alpha-galactosidase A (AGAL-A), resulting in the accumulation of glycosphingolipids, mainly globotriaosylceramide (Gb3) and its deacetylated form globotriaosylsphingosine (lysoGb3), the latter being commonly used as a surrogate biomarker [1–3]. FD has historically been characterized by acral pain, angiokeratoma, cerebral strokes, progressive renal failure and cardiomyopathy, with a diminished life-expectancy [4]. However, the clinical presentation and incidence of FD are changing as the diagnostic approach is moving from clinicobiochemical algorithms to genetic screenings. Indeed, the first estimations based on clinical ascertainment before 2000 evaluated the incidence of FD between 1:40,000–117,000 live births [5,6], whereas three recent newborn screening studies observed incidences greater than 1:10,000 [7–10]. Since 1990, a nonclassical or late-onset phenotype of FD has been described, with higher residual AGAL-A activity and predominant, if not isolated, cardiac manifestations [11]. The majority of the individuals detected by genetic screenings carry galactosidase A alpha (*GLA*) variants that are usually associated with this nonclassical phenotype of FD [7,12]. The classical and nonclassical phenotypes have been empirically determined

on the basis of the presence or absence of characteristic symptoms (usually neuropathic pain, angiokeratoma, and cornea verticillata (CV)), *GLA* enzyme activity and/or the *GLA* genetic variant, though without any consensus [12,13]. As the prognosis of the different phenotypes is markedly different, there is a need to determine reproducible classification criteria to improve the reliability of therapeutic studies and to personalize the bedside management of FD. Some scoring systems already exist, and they have been elaborated with empirical considerations; these scoring systems include many nonobjective criteria with several items that make them difficult to use in a daily practice. Moreover, existing scoring systems do not differentiate non-classical from classical phenotypes of the disease whereas a growing literature suggests the need for personalized management [14–16]. In this study, we employed unsupervised multivariate statistics for clinical data to identify simple and objective criteria that would allow an effective classification of adult patients. Additionally, we propose a new and simple scoring system based on this classification to assess the clinical severity and facilitate the management of FD patients.

Materials and methods

Patients, clinical data and biological samples

We analyzed data from patients prospectively included in the multicenter cohort FFABRY with an enzymatic and/or genetic diagnosis of FD from December 2014 to May 2017. Written consent were obtained after written and verbal information. The present study was approved by the local ethics committee (Comité de Protection des Personnes VI—Pitié Salpêtrière) and the Comité consultatif sur le traitement de l'information en matière de recherche dans le domaine de la santé, according to the relevant French legislation. Clinical data were prospectively collected through a standardized online form. Cardiac hypertrophy was defined as diastolic interventricular septum thickness > 13 mm by cardiac echocardiography or magnetic resonance imaging (MRI). Arrhythmia was defined as the presence of cardiac conduction defect or rhythm trouble. Estimation of the glomerular filtration rate (eGFR) was based on the CKD-EPI equation [17]. Glomerular hyperfiltration was defined as eGFR > 135ml/min/ 1.73m² [18]. Proteinuria was positive if above 0.3 g/24 h or if the proteinuria/creatininuria ratio was > 50 mg/ mmol. Cornea verticillata was assessed via slit-lamp examination. If not mentioned in the medical records, the patients were considered to not have a history of the following items: cerebral stroke, movement disorder, seizure, renal or cardiac transplantation, dialysis, need of a pacemaker (PM), and cardiac failure. All other items were considered missing if not mentioned. The Mainz Severity Score Index (MSSI) was calculated automatically according to the scoring system established by Whybra et al. [19].

Blood samples were collected at the time of inclusion. Plasma was isolated by centrifugation using BD Vacutainer^{∞} serum tubes with an increased silica act clot activator and BD Vacutainer^{∞} heparin tubes before storage at -80°C. All patients were screened for the presence of antiagalsidase antibodies, as previously described [20]. LysoGb3 concentrations were measured in available plasma samples (n = 36) by ultra-performance liquid chromatography coupled to tandem mass spectrometry (UPLC-MS/MS), as previously described [20].

Statistical analyses

Males and females were analyzed separately due to the known phenotype differences [12]. We performed ascending hierarchical clustering on principal components (HCPC) after multiple correspondence analysis (MCA) for the following categorical variables: presence or history of CV, angiokeratoma, history of Fabry acral pain, hypertrophic cardiomyopathy (HCM), arrhythmia, eGFR </>45 ml/min/1.73 m², renal transplant, ischemic stroke and hearing loss

and GLA variant type (missense vs. others). All the categorical variables were used as active and included in the clinical clustering except the GLA variant type used as illustrative. Cerebral MRI abnormalities were excluded due to missing data. Age was considered an illustrative variable and was not included in the clustering. MCA and HCPC were performed with R software version 3.4.0 and the package FactoMineR. Patients with missing data were excluded from this analysis. The correlation between variable and dimension was considered significant at p < 0.02. We defined the best algorithm to meet the previous clusters using ROC curves. After verification for normal distribution and equality of variances with Shapiro-Wilk and Levene tests, respectively, we employed parametric tests such as the t-test with and without Welch's correction for unequal variances, Pearson correlation and linear regression for Gaussian values, or nonparametric tests such as Kruskal-Wallis (KW) and Mann-Whitney (MW) comparison tests and the Spearman correlation test. We used the log-rank test with Kaplan-Meier to analyze the survival distribution. We applied logistic regression with stepwise selection based on p-values for discrete variables and Fisher's exact t-test for contingency. The p-value for the alpha-risk in all tests, except for MCA and HCPC, was 0.05. GraphPad Prism 5.0 and the EZR plugin version 1.35v [21] packages for R software were used.

Results

From December 2014 to May 2017, 104 patients (54 males) were prospectively included in the FFABRY cohort. Their general characteristics are described in Table 1.

Clinical clustering in males

Multiple component analysis (MCA) and hierarchical clustering on principal components (HCPC) were performed with data for 41 male patients who had available complete data (Fig 1). Their mean age was 44.4 years-old. We assume that treatment with ERT or chaperone therapy does not modify the overall phenotype of patients. Six patients were untreated at the inclusion (mean age 34.6 years-old; min-max 17.1-58.1 y.). Mean duration of treatment was 6.5 years in treated patients (min-max: 0.26-15.5 y.). The first 2 dimensions of MCA expressed 53.7% of the total inertia. An ascending HCPC performed on the first 5 dimensions identified 3 different clusters (Fig 2). Group 1 (mean age = 50.5 +/- 11.2 years; n = 21) was characterized by the absence of CV ($p < 10^{-7}$), the absence of angiokeratoma ($p < 10^{-4}$), the absence of acral pain (p<0.001), a missense mutation (p<0.02), eGFR > 45 ml/min/1.73 m² (p<0.02), and the absence of renal transplant (p = 0.02). Group 2 (mean age = 32.3 + /-9.9 years; n = 13) was characterized by the absence of hypertrophic cardiomyopathy (HCM) (p $<10^{-3}$), the presence of CV (p = 0.001), the presence of angiokeratoma (p = 0.003), the presence of acral pain (p = 0.007), and eGFR > 45 ml/min/1.73 m². Group 3 (mean age = 48.8 + /-9.5 years; n = 7) was characterized by eGFR < 45 ml/min/1.73 m² (p $< 10^{-6}$), a history of renal transplant (p $< 10^{-4}$) and the presence of CV (p = 0.002). On the basis of these characteristics, we considered group 1 to be the nonclassical phenotype and groups 2 and 3 to be the classical phenotype, with younger and older patients, respectively. When considering the absence of acral pain or CV as criteria for the nonclassical phenotype, we met the previously defined clusters with a sensitivity of 89.5%, a specificity of 91.0%, a positive predictive value of 89.5%, a negative predictive value of 91.0%, and an area under the receiver operating characteristic (ROC) curve of 0.9.

Clinical clustering in females

We applied the same approach for females using the same variables for patients with complete data (n = 36). MCA and HCPC revealed 3 different clusters without any clinical significance (Fig 3). Therefore, we considered that clinical clustering was not appropriate for females.

Table 1. Characteristics of patients (*time under enzyme replacement therapy included).

	Males (n = 54)					Females (n = 50)				
Exposure to treatment	None	Agalsidase alpha	Agalsidase beta	Agalsidase alpha and beta	Migalastat (+/- agalsidase)	None	Agalsidase alpha	Agalsidase beta	Agalsidase alpha and beta	Migalastat (+/- agalsidase)
Exposed (n)	10	12	18	11	3	25	9	8	6	2
Currently treated (%)	-	44 (81%)			-	25 (50%)				
Median age (Q1Q3)	27.2 years (20.8– 41.0)	46.1 years (34.3–53.6)	48.7 years (43.4–60.4)	42.7 years (31.9–47.8)	43.3 years (32.7–52.1)	43.2 years (36.1– 53.2)	52,7 years (48,0–54,6)	58,5 years (47,6–63,0)	54,9 years (49,5–61,2)	52.4 years +/- 8.2 years
Median follow-up time (Q1Q3)	4.1 months (1.9–5.4)	5.7 years (3.7–7.7)	7.1 years (2.8–13.8)	10.7 years (5.2–16.2)	4.2 years* (3.9–8.7)	3.0 years (0.6– 5.5)	5.8 years (3.0–12.2)	14.8 years (11.6–15.9)	8.6 years (6.2–10.3)	9.8 years* +/- 8.1 years
Median cumulative exposure to specific treatment (Q1Q3)	-	4.0 years (1.3–6.4)	4.6 years (0.3–9.3)	10.6 years (4.0–12.7)	4.2 years (0.6–8.2)*	-	2.5 years (2.4–9.9)	6.7 years (2.2–11.4)	6.4 years (5.2–10.0)	2.4 years* +/- 1.6 years
Mean age at visit (mean +/- SD)	43.4 +/- 14.7 years					48.9 +/- 14.5 years				
Mean +/- SD cumulative exposure to specific treatment	7.0 +/- 4.8 years					6.1 +/- 4.5 years				
Median MSSI Neurological [Q1-Q3] (max 20)	5 [1.3–8]					4.5 [1.3–6.0]				
MSSI Cardiac (max 20)	6 [0.0–11.8]					2 [0.0–10.8]				
MSSI Renal (max 18)	0 [0-8]				0 [0-8]					
MSSI General (max 18)	4 [2-6]				2 [1–4]					
MSSI Global (max 76)	20.5 [12.5–28.0]				14.5 [7.0–21.8]					

FFABRY score

As already mentioned, the morbidity of FD relies on renal, cardiac and central nervous system involvement. Hence, the prognosis of FD depends on the clinical phenotype of patients. Based on the results of the previous clustering, we introduce the first severity scoring system that takes into account the clinical phenotype of FD. The FFABRY score is therefore constructed with 4 variables: the overall clinical phenotype, the kidney disease score, the heart disease score and the central nervous system score as followed:

Phenotype

- Males with the classical phenotype: past or actual acroparesthesia and cornea verticillata
- Males with the nonclassical phenotype: no past or actual acroparesthesia or no cornea verticillata
- Females

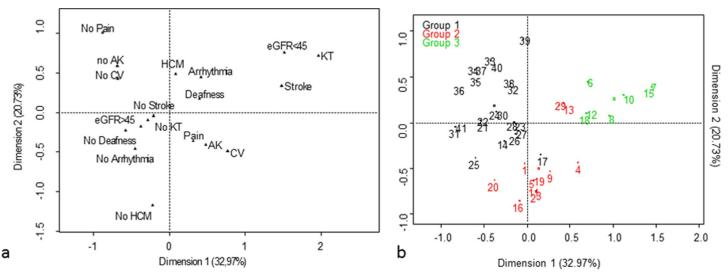


Fig 1. a: Variable factor map obtained by multiple component analysis using data for 41 males with complete data (AK: angiokeratoma; CV: cornea verticillata; eGFR: estimated glomerular filtration rate in ml/min/1.73 m²; HCM: hypertrophic cardiomyopathy; KT kidney transplantation). b: Ascending hierarchical classification of individuals using the first 5 dimensions of multiple component analysis performed using data for 41 males with complete data. Three clusters were identified. Group 1, characterized by the absence of cornea verticillata ($p < 10^{-7}$), the absence of angiokeratoma ($p < 10^{-4}$), the absence of acral pain (p < 0.001) and the absence of renal disease, was referred to as the nonclassical cluster. Groups 2 and 3, characterized by the presence of cornea verticillata (p < 0.002), were referred to as classical groups with younger (mean age 32.3 +/- 9.9 years) and older (mean age 48.8 +/- 9.5 years) patients, respectively.

Kidney disease: K score from K0 to K5. FD renal involvement is progressive and characterized by glomerular hyperfiltration and proteinuria followed by a decrease in glomerular function [22,23]. We propose the scale as follows:

K0: No proteinuria and $90 > eGFR > 135 \text{ ml/min/}1.73 \text{ m}^2$

K1: Hyperfiltration such as eGFR ≥ 135 ml/min/1.73m² without proteinuria

K2: $60 < eGFR < 90 \text{ ml/min}/1.73 \text{ m}^2 \text{ OR proteinuria} > 0.3g/24h \text{ or } 50 \text{ mg/mmol}$

K3: $30 < eGFR \le 60 \text{ ml/min}/1.73 \text{ m}^2 +/-\text{proteinuria}$.

K4: $15 < eGFR \le 30 \text{ ml/min}/1.73 \text{ m}^2 +/-\text{ proteinuria}$.

K5: $eGFR < 15 \text{ ml/min}/1.73 \text{ m}^2$ or dialysis or renal transplant.

Heart disease: H score from H0 to H4. FD is a cause of HCM, progressively leading to diastolic dysfunction, ischemia or obstructive cardiac failure [24]. FD is also characterized by arrhythmia, which has become the leading cause of death [25,26]. An interventricular septum thickness (IST) > 30 mm has been associated with a high risk for sudden death [24]. Additionally, we propose the following staging:

H0: No HCM. No cardiac symptomatology.

H1: HCM such as $13 < IST \le 30$ mm and/or QRS interval on ECG ≥ 200 msec and/or ventricular hypertrophy on ECG without cardiac symptoms and no need of antiarrhythmic or beta-blockers.

H2: H1 + need of antiarrhythmic or beta-blockers*.

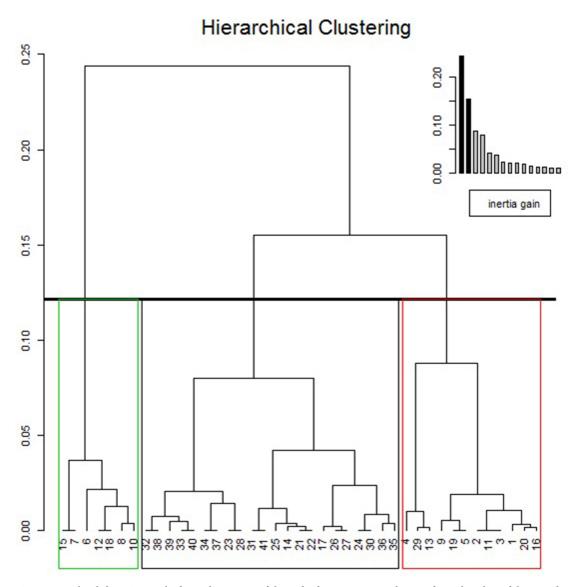


Fig 2. Hierarchical clustering on the first 5 dimensions of the multiple component analysis performed on data of the 41 males with complete data.

H3: 35% < Left ventricular ejection function (LVEF) \leq 50% and/or a need of pacemaker (PM) implantation and/or angina.

H4: Need of heart transplant or LVEF \leq 35% or IST > 30 mm.

* according to guidelines on management of HCM [27]

Central nervous system involvement: N score from N0 to N2. In a logistic regression model, we observed that chronic headaches were associated with a higher risk of stroke in males (OR 16.2, p = 0.01), independent of renal and cardiac diseases. Moreover, a trend toward a higher risk of cerebral stroke was observed in males with cochlear disorder defined by the presence of tinnitus or hearing loss (OR 7.8, p = 0.054), independent of age. Additionally, we propose the following staging:

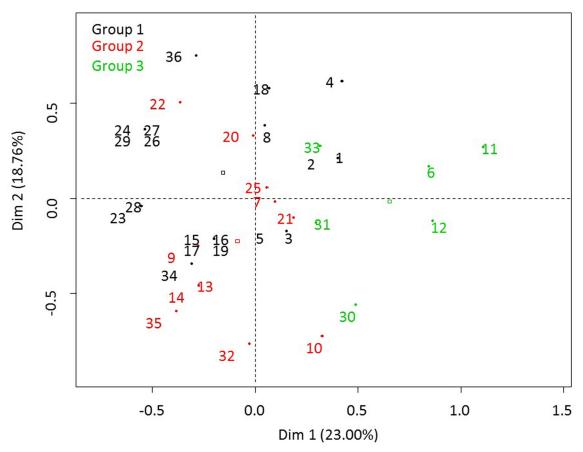


Fig 3. Ascending hierarchical classification of individuals (using the first 5 dimensions of multiple component analysis performed using data for 36 females with complete data) did not reveal any significant groups.

N0: No chronic headache and no cochlear disorder and no history of stroke.

N1: Chronic headache and/or tinnitus and/or hearing loss and no history of stroke.

N2: History of cerebral stroke.

Total FFABRY score from T0 to T11. The total (T) score was defined as the sum of the K-, H- and N-scores.

Description of the FFABRY cohort clustered by sex, cornea verticillata and acroparesthesia

Using our clustering criteria, we distinguished 29 males with the classical phenotype, 25 males with the nonclassical phenotype and 50 females. Their clinical and biochemical characteristics are described extensively in Table 2. Briefly, among males, those with the nonclassical phenotype were diagnosed at an older age (45.4 vs. 28.6 years; $p < 10^{-3}$) and had a milder phenotype including a less steep eGFR slope (-1.7 vs. -2.4 ml/min/1.72m²/ years; p < 0.05), a lower risk of renal transplantation (log-rank, HR = 0.07; $p < 10^{-3}$) and a lower risk of cerebral stroke (1/25 vs. 4/29; non-significant). Surprisingly, ENT involvement was frequent in both male groups, being observed in 70.0% of classical and 52.0% of nonclassical cases (p = ns). Of note, one of the most commonly reported symptoms was anxiety, which was observed in 70.3% of the male

Table 2. Characteristics of patients stratified by clinical phenotype defined with the FFABRY scoring system.

Phenotype	Classical Males	Nonclassical Males	Females
n	29	25	50
Median age [IQR] (years)	39.8 [29.5-46.3]	51.8 [43.0-60.8]	51 [38.0-58.5]
Median age at diagnosis (years)	28.6	45.4	42.8
Treatment received (N =)	25	19	25
Agalsidase alpha (n =)	8	4	9
Agalsidase beta (n =)	8	10	8
Both agalsidases successively (n =)	8	3	6
Migalastat (n =)	1	2	2
Mean cumulative duration of treatment (+/- SD; years)	10.3 +/- 8.5	13.1 +/- 5.1	6.1 +/- 4.0
enal disease			
R-score > 0	15	19	24
Median R-score	1	0.5	0
Median renal MSSI (IQR)	4 (0-18)	0 (0-8)	
eGFR slope in ml/min/1.73 m²/y (r²; p)	-2.4 (0.66; p < 0.0001)	-1.7 (0.61; p < 0.0001)	-0.7 (0.19; p < 0.003)
ESRD (n)	8 ¹	0	1
Median survival without renal transplantation	48.6 years	NA	NA
ACE blocker (n)	14	14	16
ACE blockers in patients with R-score >0		9%	50%
ardio-vascular disease			
HCM ² (n =)	15	213	19
Median survival without HCM	46.3 years	57.6 years ⁴	62.0 years
Pacemaker (n =)	1	6 ⁵	1
Median survival without severe cardiac event (H-score ≥ 3)	50.8 years	60.8 years ⁶	71.2 years
leurological disease	7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	2212 / 2022	, -12 , -13
Dyshidrosis (n)	18/29	6/25 (p = 0.005)	34/50
Heat intolerance (n)	17/23	8/23 (p = 0.008)	9/40
Ischemic strokes (n; %)	4; 13.8%	1; 4.0%	9; 18.0%
Median age [IQR]; years	27.6 [15.0–33.4]	34.0	45.1 [45.1–61.9]
NT involvement	27.0 [13.0-33.4]	31.0	45.1 [45.1-01.7]
Tinnitus	12/22 (54.5%)	7/22 (31.8%)	15/44 (34.1%)
Hearing loss	17/24 (70.8%)	13/25 (52.0%)	20/48 (41.7%)
Ophthalmological involvement	17/24 (70.870)	13/23 (32.0%)	20/48 (41.770)
Cataract (median age at diagnosis; years)	5/12 (44.5)	4/20 (64.3)	6/42 (60.1)
Cornea verticillata	19/19	0/23	28/42
	19/19	0/23	20/42
Other Angiokeratoma	23 (79.3%)	8 (33.3%) ⁷	10/47 (20 20/)
Angiokeratoma Abdominal pain	14/27 (51.8%)	5/17 (29.4%)	18/47 (38.3%) 8/42 (24.4%)
Abdominai pain Mental health	14/2/ (31.0%)	3/17 (29.4%)	0/42 (24.4%)
Anxiety	10/20	(70.204)	16/34 (47.1%)
Anxiety Depression symptoms	12/38 (, ,	
	11/38 (15/44 (44.1%)	
Suicide attempt		38	0
Mutations	c.137A>G*	c.334C>T*	c.123del*
	c.169C>T	c.337T>C*	c.125T>G*
	c.233C>G	c.486G>C*	c.214del (n = 2)*
	c.334C>T (n = 3)*	c.522T>A*	c.233C>G (n = 2)
	c.424T>C	$c.644A>G (n = 9)^*$	$c.334C>T (n = 5)^*$
	c.486G>C	c.692A>G*	c.424T>C*

(Continued)

Table 2. (Continued)

	c.539del*	$c.713G>A(n=2)^*$	c.427G>A (n = 2)*	
	c.548G>C*	c.758T>C	c.486G>C*	
	c.680G>A	c.802-3_802-2del*	c.504A>C*	
	c.729G>C*	c.847C>T (n = 2)*	c.548G>C*	
	$c.798T > A (n = 2)^*$	c.902G>A*	c.655A>C*	
	c.802-3_802-2del*	c.1010T>C*	c.680G>A*	
	c.806T>C*	c.1016T>G*	c.695T>C*	
	c.847C>T*	$c.1087C>T (n = 2)^*$	c.718_719del*	
	c.875C>T		c.729G>C*	
	c.884T>G		c.798T>A	
	c.901C>T (n = 2)*		c.802-3_802-2del (n = 4)*	
	c.902G>A*		c.840A>T	
	c.1010T>C*		c.884T>G (n = 3)*	
	c.1069_1079del*		$c.901C>T (n = 6)^*$	
	c.1246C>T*		c.902G>A*	
	no data (n = 4)		c.1277_1278del (n = 2)*	
			c.1021del	
			c.1024C>T	
			c.1087C>T	
			c.1168G>A	
			no data (n = 6)	
ysoGb3 (median in ng/ml; IQR; n)				
In treated patients	18.9 (11.6–32.3; n = 17)	$6.25 (2.6-21.9; n = 17)^8$	4.5 (2.7–6.2; n = 15)	
In untreated patients	101.8 (n = 2)	8.5 (3.0–16.7; n = 5)	2.6 (1.7–3.8; n = 22)	

eGFR: estimated glomerular filtration rate; ESRD: end-stage renal disease; ACE-blocker: angiotensin conversion enzyme blocker; HCM: hypertrophic cardiomyopathy; PM: pacemaker; NA: not available. (1) HR vs. nonclassical 14.4; log-rank p=0.0003; (2) in Cox regression, HCM is influenced by age at diagnosis (HR 0.81; p<10-7) and cumulative exposure to treatment (HR 0.71; p<0.0001); (3) in Cox regression, influenced by the phenotype: HR nonclassical vs. classic: 0.19; p=0.006); (4) log-rank classical vs. nonclassical; p<0.003; (5) log-rank classical vs. nonclassical; p=0.01; (7) Fisher-exact t-test classical vs. nonclassical; p<0.001; (8) Mann-Whitney, p=0.01

*variants included in the MCA and HCPC analyses (exhaustive data). One 46.6-year-old female, a p.A143T carrier, was included in the MCA and HCPC analyses. She had been treated for 2 years with migalastat for acral and abdominal pain and angiokeratoma but had no renal, cardiac or cerebrovascular involvement (MSSI = 13; plasma lysoGb3 1.1nM). Another female p.A143T carrier, aged 43.0 years, had no dermatological, ophthalmological, neurological, renal or cardiological symptoms (MSSI = 1; plasma lysoGb3 1.3nM). She received no specific treatment and was not included in the MCA and HCPC analyses due to missing data.

https://doi.org/10.1371/journal.pone.0233460.t002

patients, with suicide attempts reported for 10.5% of the male cohort. Females exhibited a contrasting globally milder phenotype (Fig 4 and Table 2). However, 18.0% of them had a history of ischemic stroke, with a median age at diagnosis of 45.1 years, and the event was significantly associated with the existence of cardiac rhythm problems (OR: 15.3; p = 0.046). Such an association with rhythm problems was not observed in men.

Performances of clustering with MSSI and FFABRY scores in the entire FFABRY cohort

Clustering appeared clinically relevant using both MSSI and FFABRY scores, with significant differences between groups stratified by classes of ages (Table 3). The classical phenotype was associated with a higher risk of severe renal events (K-score \geq 3; Cox analysis: HR (classical/nonclassical) = 35.1; p <10⁻³; HR (classical/females) = 100; p < 10⁻⁵; Fig 4) and a higher risk of severe cardiac events (H-score \geq 3; Cox analysis: HR (classical vs. nonclassical) = 4.8;

Severe event-free survival

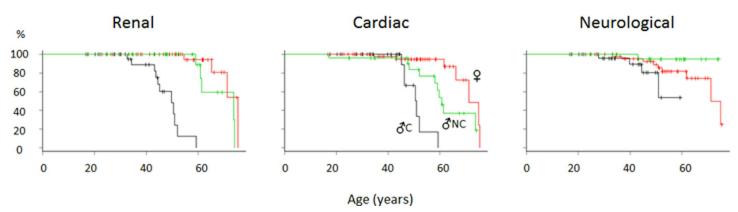


Fig 4. Renal, cardiac and neurological severe event-free survival curves (black: Classical group males, green: Nonclassical group males, red: Females).

https://doi.org/10.1371/journal.pone.0233460.g004

p = 0.008; HR (classical/ females) = 16.7; $p < 10^{-4}$; Fig 4), and there was a trend toward a higher risk of severe neurological events (N-score = 2) in males with the classical compared to the nonclassical phenotype (Cox analysis; HR = 7.7; p = 0.08, Fig 4). There was no significant difference among females (HR = 2.5; p = 0.2). T-score evolution illustrated that the overall severity increased with time in the classical group compared to the nonclassical group (Fig 5).

LysoGb3 plasma levels, anti-agalsidase antibodies and FFABRY

LysoGb3 plasma levels were higher in males with the classical phenotype compared to those with the nonclassical phenotype or females in both treatment-naïve (respective medians 101.8, 5.8 and 3.2; Kruskal Wallis p < 0.04) and in treated (respective medians 18.9, 6.7 and 4.5; $p < 10^{-4}$) patients. As expected, there was no correlation between lysoGb3 plasma levels and FFABRY scores after stratification by phenotype among treated patients. Two female and 18 male patients were positive for anti-agalsidase antibodies. If considering only males who had been exposed to agalsidase, the presence of antibodies was significantly associated with the classical phenotype (56% vs. 20%, p < 0.02). Their characteristics have already been reported [20].

Genotype-phenotype correlation

Among the 30 different genetic variants observed in males, nonsense variants were associated with the classical phenotype (9/13 patients with nonsense variants). The four patients classified in the nonclassical group had no cornea verticillata: a c.802-3_802-2del carrier with a K1H3N0 score at 17.1 years, two c.847C>T carriers with K0H1N0 and K0H1N1 scores at 33.4 and 47.3 years, and a c.522T>A carrier with a K2H0N0 score at 47.7 years. Five variants were observed in both the classical and nonclassical groups: c.334C>T, c.847C>T, c.802-3_802-2del, c. 902G>A and c.1010T>C. Their characteristics are described in Table 4.

Discussion

The nonclassical phenotype of FD has become the most prevalent [7,12]. However, most clinical and fundamental research studies addressing with FD have not stratified patients based on phenotype. Here, we propose a simple algorithm to distinguish patients. We demonstrate that CV assessed by slit-lamp and acral pain, but not angiokeratoma, are statistically sufficient to

 $Table \ 3. \ FFABRY \ and \ MSSI \ scores \ (comparison \ with \ Mann-Whitney \ test \ p^*: Classical \ versus \ nonclassical \ males; \ p^{**}: Males \ versus \ females).$

	Males			Females	
	Classical	Nonclassical	p *		p **
n	29	25		50	
Age total	39.84 [29.49, 46.29]	51.76 [42.96, 60.76]	0.001	51.00 [37.99, 58.51]	0.001
< 35 y n (median [IQR])	14 (28.69 [21.69, 32.51])	6 (29.20 [24.90, 34.35])	0.509	10 (27.16 [25.09, 32.51])	0.766
35-50 y n (median [IQR])	11 (44.50 [43.52, 46.30])	6 (47.46 [44.28, 47.66])	0.228	13 (43.14 [38.39, 46.56])	0.233
> 50 y n (median [IQR])	4 (51.58 [50.93, 53.93])	13 (60.76 [58.10, 67.58])	0.024	27 (58.22 [52.95, 65.26])	0.074
FFABRY Heart score	1.00 [0.00, 2.00]	2.00 [1.00, 3.00]	0.085	1.00 [0.00, 2.00]	0.025
< 35 y (median [IQR])	0.00 [0.00, 1.00]	1.00 [0.25, 1.75]	0.130	0.00 [0.00, 0.00]	0.213
35–50 y (median [IQR])	1.00 [1.00, 2.50]	1.50 [0.25, 2.75]	0.716	0.00 [0.00, 2.00]	0.113
> 50 y (median [IQR])	3.50 [3.00, 4.00]	3.00 [2.00, 3.00]	0.040	2.00 [1.00, 2.00]	0.001
FFABRY Kidney score	1.00 [0.00, 5.00]	0.50 [0.00, 2.00]	0.214	0.00 [0.00, 2.00]	0.228
< 35 y (median [IQR])	0.00 [0.00, 0.75]	0.00 [0.00, 0.75]	1.000	0.00 [0.00, 0.00]	0.902
35–50 y (median [IQR])	2.00 [1.00, 5.00]	0.00 [0.00, 0.00]	0.024	1.00 [0.00, 2.00]	0.026
> 50 y (median [IQR])	4.50 [4.00, 5.00]	2.00 [1.50, 3.00]	0.004	2.00 [0.00, 2.00]	0.004
FFABRY Neurological score	1.00 [0.00, 1.00]	1.00 [0.00, 1.00]	0.340	1.00 [0.00, 1.00]	0.669
< 35 y (median [IQR])	0.50 [0.00, 1.00]	0.00 [0.00, 0.00]	0.165	1.00 [0.00, 1.00]	0.231
35–50 y (median [IQR])	1.00 [0.25, 1.00]	1.00 [0.25, 1.00]	0.859	0.00 [0.00, 1.00]	0.743
> 50 y (median [IQR])	1.00 [1.00, 1.25]	1.00 [0.00, 1.00]	0.076	1.00 [0.00, 1.00]	0.311
FFABRY Total score	3.00 [1.00, 7.00]	4.00 [2.00, 6.00]	0.882	2.50 [1.00, 4.75]	0.312
< 35 y (median [IQR])	1.00 [1.00, 1.75]	1.50 [1.00, 2.00]	0.724	1.00 [0.00, 1.00]	0.487
35–50 y (median [IQR])	5.50 [3.25, 7.00]	2.50 [2.00, 3.75]	0.043	2.00 [1.00, 4.25]	0.030
> 50 y (median [IQR])	9.00 [8.00, 10.25]	6.00 [4.00, 6.00]	0.003	3.00 [2.00, 5.00]	0.002
MSSI cardiac	2.00 [0.00, 9.00]	10.00 [3.00, 14.00]	0.013	2.00 [0.00, 10.75]	0.025
< 35 y (median [IQR])	0.50 [0.00, 2.75]	4.50 [0.00, 9.00]	0.505	0.00 [0.00, 1.50]	0.348
35–50 y (median [IQR])	6.00 [2.00, 10.50]	6.50 [1.50, 9.25]	0.724	0.00 [0.00, 2.00]	0.047
> 50 y (median [IQR])	1.50 [0.00, 6.75]	14.00 [13.00, 16.00]	0.190	9.00 [1.50, 14.00]	0.067
MSSI general	5.00 [3.00, 8.00]	2.00 [1.00, 4.00]	0.001	2.00 [1.00, 4.00]	< 0.001
< 35 y (median [IQR])	4.50 [2.00, 6.00]	2.00 [1.25, 2.00]	0.054	1.50 [1.00, 4.75]	0.167
35–50 y (median [IQR])	5.00 [4.00, 8.50]	5.50 [3.25, 7.00]	0.577	1.00 [1.00, 3.00]	0.005
> 50 y (median [IQR])	6.50 [5.00, 9.25]	1.00 [1.00, 2.00]	0.006	2.00 [1.50, 4.00]	0.006
MSSI renal	4.00 [0.00, 18.00]	0.00 [0.00, 8.00]	0.052	0.00 [0.00, 8.00]	0.111
< 35 y (median [IQR])	0.00 [0.00, 0.00]	0.00 [0.00, 0.00]	0.342	0.00 [0.00, 0.00]	0.558
35–50 y (median [IQR])	8.00 [6.00, 18.00]	0.00 [0.00, 0.00]	0.015	4.00 [0.00, 4.00]	0.010
> 50 y (median [IQR])	13.00 [8.00, 18.00]	8.00 [0.00, 8.00]	0.017	0.00 [0.00, 8.00]	0.030
MSSI neurological	7.00 [3.00, 10.00]	2.00 [0.00, 5.00]	0.001	4.50 [1.25, 6.00]	0.002
< 35 y (median [IQR])	6.00 [2.25, 8.00]	4.00 [2.25, 5.75]	0.534	5.50 [5.00, 7.50]	0.640
35–50 y (median [IQR])	6.00 [3.50, 12.00]	4.00 [0.75, 7.25]	0.362	5.00 [1.00, 6.00]	0.348
> 50 y (median [IQR])	9.00 [7.75, 10.75]	1.00 [0.00, 2.00]	0.004	3.00 [0.00, 5.50]	0.004
MSSI total	24.00 [14.00, 33.00]	20.00 [12.00, 24.00]	0.080	14.50 [7.00, 21.75]	0.017
< 35 y (median [IQR])	14.00 [8.75, 18.75]	10.50 [4.75, 18.50]	0.535	11.00 [5.50, 18.50]	0.704
35–50 y (median [IQR])	29.00 [24.50, 32.50]	16.50 [12.00, 24.75]	0.039	13.00 [7.00, 15.00]	0.002
> 50 y (median [IQR])	35.50 [33.75, 37.25]	22.00 [18.00, 24.00]	0.005	19.00 [9.00, 26.00]	0.014

distinguish males with the classical phenotype from those with the nonclassical phenotype, with significant differences in terms of severity. As expected, males with the classical phenotype have more severe renal disease than do males with the nonclassical phenotype, but the

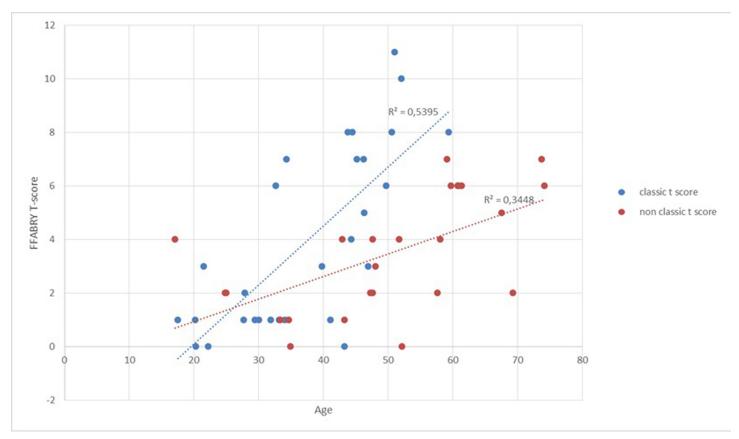


Fig 5. T-score evolution in the classical group compared to the nonclassical group (with associated linear regression curve).

former also experience cardiomyopathy earlier, and some of them also have stroke early, independent of arrhythmia. Females cannot be separated into classical and nonclassical phenotypes, likely due to the X-inactivation status in the different organs.

Hemangiomas, which are frequent, can easily be misdiagnosed as angiokeratoma, which may explain the reduced specificity of angiokeratoma for the classical phenotype. In contrast, CV is a pivotal criterion in our classification. Although CV can be related to exposure to amiodarone, its association with the severity of FD has already been described, as observed in 94% of classical Fabry cases [28]. A recent paper reported that CV is often underdiagnosed in FD

Table 4. Characteristics of patients with genetic variants observed in both classical and nonclassical groups (eGFR: Estimated glomerular filtration by CKD-EPI equation in ml/min/1.73m²; RT: Renal transplant).

Variant	Classical patient							
	age	Treatment duration (y)	FFABRY score	eGFR	age	Treatment duration (y)	FFABRY score	eGFR
c.334C>T	34.4	6.2	K5 H1 N1 T7	RT	48.1	13.4	K0 H3 N0 T3	105
	45.2	3.4	K3 H3 N1 T7	52				
	59.4	5.6	K4 H3 N1 T8	23				
c.802-3_802-2del	50.6	1.3	K4 H3 N1 T8	25	17.1	0	K1 H3 N0 T4	136
c.847C>T	49.8	11.1	K5 H1 N0 T6	RT	33.4	13.3	K0 H1 N0 T1	116
					47.3	13	K0 H1 N1 T2	102
c.902G>A	17.5	0.4	K1 H0 N0 T1	136	24.9	0	K0 H2 N0 T2	121
c.1010T>C	46.3	4.3	K2 H3 N0 T5	81	60.9	4.8	K3 H2 N1 T6	57

https://doi.org/10.1371/journal.pone.0233460.t004

using the slit-lamp approach [29]. The present work concerned 4 males whose phenotype was not mentioned and 10 females. However, whether a more sensitive approach using in vivo corneal confocal microscopy (IVCM) may reveal minimal deposits in nonclassical patients and females is not the question. Hence, we suggest that an unquestionable CV, with obvious deposits assessed by slit-lamp, is associated with a classical FD phenotype. Similarly, our second pivotal criterion is the existence past or actual of acral pain and not the proof of small fiber neuropathy assessed by paraclinical exams.

Our study was based on data for 104 adult patients; a small number of patients is inherent with the rarity of FD. However, FFABRY is a multicenter database including patients from different pedigrees, different medical specialties (nephrology, cardiology, internal medicine, and genetics) and different locations in France, which allows for clinical and genetic heterogeneity (42 different variants) that benefits analyses. We used a linear regression model to assess the slope of eGFR evolution in the cohort, which is limited by the heterogeneity of patients, especially females. Analysis of X-inactivation may allow for a better classification of the female phenotype and help in stratifying the risk of severe events; unfortunately, this analysis remains unavailable in routine practice [30]. As experts managing lysosomal diseases in a dedicated tertiary center, we assume that treatment with ERT or chaperone therapy has not modified the overall natural history of the disease or the prognosis of patients. Nonetheless, to the best of our knowledge, regression but no disappearance of CV has ever been reported [31].

In the era of evidence-based medicine, severity scores have become mandatory for evaluating therapeutics. With FFABRY, we propose a clinically and statistically objective severity scoring system for FD. The FFABRY score was developed based on the natural history of FD and the severe clinically relevant events we observed in our center of expertise for lysosomal diseases. Our N-score highlights the importance of cochlear disorders and headaches, which are associated with the risk of cerebral stroke in males, whereas white matter lesions (WMLs) were not related to any specific symptom.

FFABRY allowed us to establish a portrait of current Fabry patients, distinguishing males with classical and nonclassical phenotypes and females with their proper clinical specificity. Interestingly, we observed no systematic genotype-phenotype correlation. Some patients sharing the same genetic variant were classified into two different groups. The FFABRY scores as well as individual clinical criteria demonstrated that our classification performed better than genotype for describing disease severity. Other scoring systems have already been developed for FD. The MSSI, which has been the most commonly used scoring system, was developed on the basis of data from 24 males and 15 females registered in the Fabry Outcome Survey (FOS) database (Shire-Takeda)[19]. The MSSI includes 26 variables, empirically weighted, among which hemorrhoids, facial appearance and subjective fitness assessment are notable; however, the MSSI does not include renal transplantation. The Disease Severity Scoring System (DS-3), which was elaborated by experts from the Fabry Registry (Sanofi-Genzyme), also includes nonobjective items such as the "patient reported domain" or sweating capacity. WMLs are included, though they are asymptomatic and not associated with poorer outcomes. Finally, the different items and their weightings have been empirically and not statistically established in these two scoring systems. Moreover, the number of items makes them challenging to use and time consuming. The Fabry International Prognostic Index (FIPI) appears to be much more robust, as it has been developed on the basis of multivariate analyses of data from 1483 patients from the FOS registry [32]. Although FIPI appears to be an effective prediction tool, it does not allow assessment of the actual clinical severity. Indeed, five of the six variables included in the cardiac item refer to extracardiac symptoms (eGFR, proteinuria, deafness, vertigo, and angiokeratoma). The online Fabry Stabilization index (FASTEX) was recently validated [33], and the authors established an attractive tool for personal follow-up of individuals.

Nevertheless, clinical phenotypes are not distinguished in this system, which could be misleading for interindividual comparisons, making the scoring system useless in group studies. After stratification according to phenotype, FFABRY scores allowed a rapid and clinically relevant evaluation of disease severity. Analyses of the K score highlight the suboptimal management of FD females in our cohort. Our study also emphasizes windows of opportunity for the diagnosis and introduction of specific treatments, as follows: before 30 years old in males with the classical phenotype, 45 years old in males with the nonclassical classical phenotype and 50 years old in females. Regarding the obviously different prognoses observed with the FFABRY score, we believe that the classical and nonclassical phenotypes should be considered as two different subtypes of FD in males, with different management strategies. As for Niemann-Pick disease A and B or Gaucher types 1, 2 and 3, it would be useful to rename the phenotypes for male patients, with classical as type 1 and nonclassical as type 2, to fully take into account such obvious clinical differences and possible pathophysiological differences.

Identifying acral pain and cornea verticillata is a rapid and simple approach that statistically discriminates Fabry phenotypes.

Supporting information

S1 Data. Clinical and biological data of included patients. (XLSX)

Acknowledgments

We gratefully thank the Société Nationale Française de Médecine Interne and Vaincre les maladies lysosomales patient association for their support as well as Isabelle Citerne and Epiconcept's team for their help.

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References

- Brady RO, Gal AE, Bradley RM, Martensson E, Warshaw AL, Laster L. Enzymatic defect in Fabry's disease. Ceramidetrihexosidase deficiency. N Engl J Med. 1967; 276: 1163–1167. https://doi.org/10.1056/NEJM196705252762101 PMID: 6023233
- Smid BE, van der Tol L, Biegstraaten M, Linthorst GE, Hollak CEM, Poorthuis BJHM. Plasma globotriaosylsphingosine in relation to phenotypes of Fabry disease. J Med Genet. 2015; 52: 262–268. https://doi.org/10.1136/jmedgenet-2014-102872 PMID: 25596309
- Young-Gqamana B, Brignol N, Chang H-H, Khanna R, Soska R, Fuller M, et al. Migalastat HCl reduces globotriaosylsphingosine (lyso-Gb3) in Fabry transgenic mice and in the plasma of Fabry patients. PloS One. 2013; 8: e57631. https://doi.org/10.1371/journal.pone.0057631 PMID: 23472096
- Schiffmann R, Warnock DG, Banikazemi M, Bultas J, Linthorst GE, Packman S, et al. Fabry disease: progression of nephropathy, and prevalence of cardiac and cerebrovascular events before enzyme replacement therapy. Nephrol Dial Transplant Off Publ Eur Dial Transpl Assoc—Eur Ren Assoc. 2009; 24: 2102–2111. https://doi.org/10.1093/ndt/gfp031 PMID: 19218538
- Meikle PJ, Hopwood JJ, Clague AE, Carey WF. Prevalence of lysosomal storage disorders. JAMA. 1999; 281: 249–254. https://doi.org/10.1001/jama.281.3.249 PMID: 9918480
- Poorthuis BJ, Wevers RA, Kleijer WJ, Groener JE, de Jong JG, van Weely S, et al. The frequency of lysosomal storage diseases in The Netherlands. Hum Genet. 1999; 105: 151–156. https://doi.org/10. 1007/s004399900075 PMID: 10480370
- Spada M, Pagliardini S, Yasuda M, Tukel T, Thiagarajan G, Sakuraba H, et al. High incidence of lateronset fabry disease revealed by newborn screening. Am J Hum Genet. 2006; 79: 31–40. https://doi.org/10.1086/504601 PMID: 16773563
- 8. Burton BK, Charrow J, Hoganson GE, Waggoner D, Tinkle B, Braddock SR, et al. Newborn Screening for Lysosomal Storage Disorders in Illinois: The Initial 15-Month Experience. J Pediatr. 2017; 190: 130–135. https://doi.org/10.1016/j.jpeds.2017.06.048 PMID: 28728811
- Hopkins PV, Klug T, Vermette L, Raburn-Miller J, Kiesling J, Rogers S. Incidence of 4 Lysosomal Storage Disorders From 4 Years of Newborn Screening. JAMA Pediatr. 2018; 172: 696–697. https://doi.org/10.1001/jamapediatrics.2018.0263 PMID: 29813145
- Wittmann J, Karg E, Turi S, Legnini E, Wittmann G, Giese A-K, et al. Newborn screening for lysosomal storage disorders in hungary. JIMD Rep. 2012; 6: 117–125. https://doi.org/10.1007/8904_2012_130 PMID: 23430949
- von Scheidt W, Eng CM, Fitzmaurice TF, Erdmann E, Hübner G, Olsen EG, et al. An atypical variant of Fabry's disease with manifestations confined to the myocardium. N Engl J Med. 1991; 324: 395–399. https://doi.org/10.1056/NEJM199102073240607 PMID: 1846223
- Arends M, Wanner C, Hughes D, Mehta A, Oder D, Watkinson OT, et al. Characterization of Classical and Nonclassical Fabry Disease: A Multicenter Study. J Am Soc Nephrol. 2017; 28: 1631–1641. https:// doi.org/10.1681/ASN.2016090964 PMID: 27979989
- 13. van der Tol L, Smid BE, Poorthuis BJHM, Biegstraaten M, Deprez RHL, Linthorst GE, et al. A systematic review on screening for Fabry disease: prevalence of individuals with genetic variants of unknown significance. J Med Genet. 2014; 51: 1–9. https://doi.org/10.1136/jmedgenet-2013-101857 PMID: 23922385
- Ortiz A, Germain DP, Desnick RJ, Politei J, Mauer M, Burlina A, et al. Fabry disease revisited: Management and treatment recommendations for adult patients. Mol Genet Metab. 2018; 123: 416–427. https://doi.org/10.1016/j.ymgme.2018.02.014 PMID: 29530533
- Citro V, Cammisa M, Liguori L, Cimmaruta C, Lukas J, Cubellis MV, et al. The Large Phenotypic Spectrum of Fabry Disease Requires Graduated Diagnosis and Personalized Therapy: A Meta-Analysis Can Help to Differentiate Missense Mutations. Int J Mol Sci. 2016; 17. https://doi.org/10.3390/ijms17122010 PMID: 27916943
- Oder D, Nordbeck P, Wanner C. Long Term Treatment with Enzyme Replacement Therapy in Patients with Fabry Disease. Nephron. 2016; 134: 30–36. https://doi.org/10.1159/000448968 PMID: 27576727

- Chen F, Du M, Blumberg JB, Chui KKH, Ruan M, Rogers G, et al. Association Between Dietary Supplement Use, Nutrient Intake, and Mortality Among US Adults: A Cohort Study. Ann Intern Med. 2019; 170: 604–613. https://doi.org/10.7326/M18-2478 PMID: 30959527
- Tonneijck L, Muskiet MHA, Smits MM, van Bommel EJ, Heerspink HJL, van Raalte DH, et al. Glomerular Hyperfiltration in Diabetes: Mechanisms, Clinical Significance, and Treatment. J Am Soc Nephrol JASN. 2017; 28: 1023–1039. https://doi.org/10.1681/ASN.2016060666 PMID: 28143897
- Whybra C, Kampmann C, Krummenauer F, Ries M, Mengel E, Miebach E, et al. The Mainz Severity Score Index: a new instrument for quantifying the Anderson-Fabry disease phenotype, and the response of patients to enzyme replacement therapy. Clin Genet. 2004; 65: 299–307. https://doi.org/ 10.1111/j.1399-0004.2004.00219.x PMID: 15025723
- 20. Mauhin W, Lidove O, Amelin D, Lamari F, Caillaud C, Mingozzi F, et al. Deep characterization of the anti-drug antibodies developed in Fabry disease patients, a prospective analysis from the French multi-center cohort FFABRY. Orphanet J Rare Dis. 2018; 13: 127. https://doi.org/10.1186/s13023-018-0877-4 PMID: 30064518
- Kanda Y. Investigation of the freely available easy-to-use software "EZR" for medical statistics. Bone Marrow Transplant. 2013; 48: 452–458. https://doi.org/10.1038/bmt.2012.244 PMID: 23208313
- Tøndel C, Kanai T, Larsen KK, Ito S, Politei JM, Warnock DG, et al. Foot process effacement is an early marker of nephropathy in young classic Fabry patients without albuminuria. Nephron. 2015; 129: 16– 21. https://doi.org/10.1159/000369309 PMID: 25531941
- Riccio E, Sabbatini M, Bruzzese D, Annicchiarico Petruzzelli L, Pellegrino A, Spinelli L, et al. Glomerular Hyperfiltration: An Early Marker of Nephropathy in Fabry Disease. Nephron. 2019; 141: 10–17. https://doi.org/10.1159/000493469 PMID: 30466100
- Marian AJ, Braunwald E. Hypertrophic Cardiomyopathy: Genetics, Pathogenesis, Clinical Manifestations, Diagnosis, and Therapy. Circ Res. 2017; 121: 749–770. https://doi.org/10.1161/CIRCRESAHA. 117.311059 PMID: 28912181
- 25. Baig S, Edward NC, Kotecha D, Liu B, Nordin S, Kozor R, et al. Ventricular arrhythmia and sudden cardiac death in Fabry disease: a systematic review of risk factors in clinical practice. Eur Eur Pacing Arrhythm Card Electrophysiol J Work Groups Card Pacing Arrhythm Card Cell Electrophysiol Eur Soc Cardiol. 2018; 20: f153–f161. https://doi.org/10.1093/europace/eux261 PMID: 29045633
- Mehta A, Clarke JTR, Giugliani R, Elliott P, Linhart A, Beck M, et al. Natural course of Fabry disease: changing pattern of causes of death in FOS—Fabry Outcome Survey. J Med Genet. 2009; 46: 548–552. https://doi.org/10.1136/jmg.2008.065904 PMID: 19473999
- 27. Authors/Task Force members, Elliott PM, Anastasakis A, Borger MA, Borggrefe M, Cecchi F, et al. 2014 ESC Guidelines on diagnosis and management of hypertrophic cardiomyopathy: the Task Force for the Diagnosis and Management of Hypertrophic Cardiomyopathy of the European Society of Cardiology (ESC). Eur Heart J. 2014; 35: 2733–2779. https://doi.org/10.1093/eurheartj/ehu284 PMID: 25173338
- 28. van der Tol L, Sminia ML, Hollak CEM, Biegstraaten M. Cornea verticillata supports a diagnosis of Fabry disease in non-classical phenotypes: results from the Dutch cohort and a systematic review. Br J Ophthalmol. 2016; 100: 3–8. https://doi.org/10.1136/bjophthalmol-2014-306433 PMID: 25677671
- Leonardi A, Carraro G, Modugno RL, Rossomando V, Scalora T, Lazzarini D, et al. Cornea verticillata in Fabry disease: a comparative study between slit-lamp examination and in vivo corneal confocal microscopy. Br J Ophthalmol. 2019. https://doi.org/10.1136/bjophthalmol-2019-314249 PMID: 31401555
- **30.** Echevarria L, Benistan K, Toussaint A, Dubourg O, Hagege AA, Eladari D, et al. X-chromosome inactivation in female patients with Fabry disease. Clin Genet. 2016; 89: 44–54. https://doi.org/10.1111/cge. 12613 PMID: 25974833
- 31. Prominent regression of corneal deposits in Fabry disease 16 years after initiation of enzyme replacement therapy.—PubMed—NCBI. [cited 27 Nov 2019]. Available: https://www.ncbi.nlm.nih.gov/pubmed/?term=Prominent+regression+ofcorneal+deposits+in+Fabrydisease+16+years+afterinitiation+of+enzymereplacement+therapy
- Hughes DA, Malmenäs M, Deegan PB, Elliott PM, Ginsberg L, Hajioff D, et al. Fabry International Prognostic Index: a predictive severity score for Anderson-Fabry disease. J Med Genet. 2012; 49: 212–220. https://doi.org/10.1136/jmedgenet-2011-100407 PMID: 22315436
- Cairns T, Wanner C. Will the FAbry STabilization indEX make its way to everyday clinical practice? Clin Kidney J. 2019; 12: 61–64. https://doi.org/10.1093/ckj/sfy126 PMID: 30747153