

Incidence and prevalence of newborn, paediatric and adolescent patients with Fabry disease: a targeted literature review

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BACKGROUND

- Fabry disease is an X-linked, multisystemic disorder caused by *GLA* gene variants resulting in α -galactosidase A (α -Gal A) deficiency.¹ Insufficient enzyme activity leads to the lysosomal accumulation of globotriaosylceramide (GL3) and other products.^{1,2}
 - These products tend to accumulate in kidney cells in utero, and continue to accumulate throughout childhood and adulthood³
 - Over time, this accumulation damages cells, leading to progressive and irreversible damage to multi-organ systems, such as the cardiovascular, renal and nervous systems.^{1,2,4-6}
- Classically affected hemizygous males, with no or very low residual α -Gal A activity, may display all of the characteristic multisystemic signs of the disease.¹
- The estimated prevalence of classic Fabry disease in newborn males has historically been estimated at 1:40,000–1:60,000.⁷
- However, newborn screening studies in Italy, Taiwan, Japan, Spain and the United States found a prevalence of *GLA* gene and α -Gal A variants in both males and females of 1:2,445–1:7,800,⁸ which is much higher than the historical estimated prevalence of classic Fabry disease.⁷
 - For newborn males reported in some of those studies, the prevalence of *GLA* gene and α -Gal A variants was as high as 1:3,100–1:7,575⁸
 - However, not all *GLA* gene and α -Gal A variants detected by screening led to symptomatic Fabry disease in all people with those variants.¹

OBJECTIVE

- To better understand the incidence and prevalence of Fabry disease and associated *GLA* gene and α -Gal A variants in newborn, paediatric and adolescent populations.

METHODS

- A literature search was conducted on data published between 1974 and 2020 (last access date: 27 August 2020) using MEDLINE/PubMed and EMBASE databases to identify epidemiology data relevant to paediatric and/or adolescent patients with Fabry disease.
 - Table 1 details the study eligibility criteria.

Table 1. Eligibility criteria for identification of relevant studies

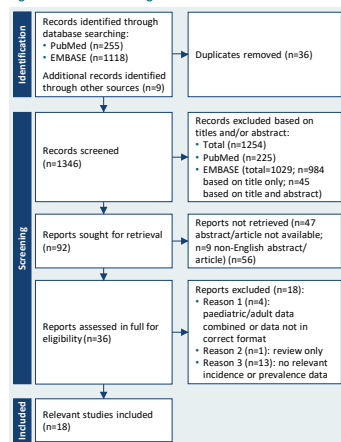
| Population | Newborn, paediatric and/or adolescent patients (aged 0–18 years) with Fabry disease |
|--------------|---|
| Intervention | Any interventions (or none) |
| Comparators | Any comparators (or none) |
| Outcomes | Includes Fabry disease incidence and/or prevalence |
| Study design | Randomised clinical trial, clinical trial, systematic review, and real-world observation studies (case, cohort or cross-sectional) |
| Other | <ul style="list-style-type: none"> Studies undertaken from 1974 to 2020 Studies published in English language Conference abstracts were eligible for inclusion |

- The search string used the terms: ‘Fabry disease’, ‘paediatric’ or ‘pediatric’, ‘adolescent’, ‘children’, ‘infant’ and/or ‘enzyme replacement therapy’ in the title and/or abstract only.

- This broad search strategy was performed to identify as many relevant articles as possible, assessed through manual screening by two independent reviewers.

- The search resulted in 1346 potentially relevant articles that underwent manual title/abstract review. Of these, 92 articles were reviewed in full for eligibility, with 18 final articles identified with relevant data (Figure 1).

Figure 1. PRISMA flow diagram



RESULTS

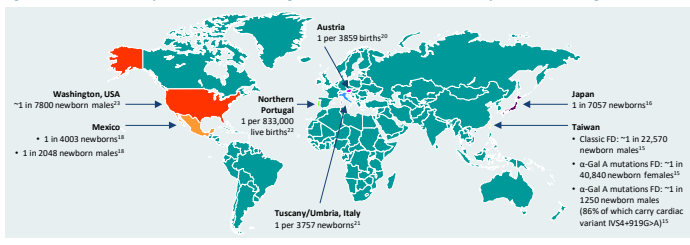
- Results are shown in Table 2.
- 18 eligible studies were included:
 - 16 studies were retrieved from EMBASE: 14 newborn screening studies with relevant incidence/prevalence data, and 2 studies that estimated prevalence based on the number of cases that had been diagnosed in a given period and the birth rate over that period.
 - 2 additional studies were retrieved from PubMed: 1 newborn screening study and 1 relevant prevalence study in an adolescent/paediatric cohort.
- The prevalence of Fabry disease *GLA* gene and α -Gal A variants in newborn screening data was found to vary by country (Figure 2).
- Overall, our prevalence estimates, based on *GLA* gene and α -Gal A variants detected in newborns, tended to be higher than historical classic male Fabry disease prevalence estimates.⁷

Table 2. Eligible studies reporting the incidence and/or prevalence of Fabry disease or associated *GLA* gene and α -Gal A variants in newborn, paediatric and/or adolescent populations

| Study | Study design | Geographical location | Population | Screening method | Prevalence | Incidence |
|---|----------------------|---------------------------|---|------------------|---|------------------------|
| Goncalves et al 2017 ⁸ | Observational cohort | Portugal | Children with juvenile idiopathic arthritis (N=292) | G | Late-onset FD-associated variant (R118C): 0.34% (1 in 292) | |
| Burlina et al 2018 ¹⁰ | Newborn screening | Italy (North East) | Newborns (N=44,411) | EA + G | | FD: 1 in 8882 |
| Burton et al 2017 ¹¹ | Newborn screening | USA (Chicago) | Newborns (N=219,793) | EA + G | | FD: 1 in 8454 |
| Chien et al 2020 ¹² | Newborn screening | Taiwan | Newborns (N=73,743) | EA + G | | FD: 1 in 18,436 |
| Colon et al 2017 ¹³ | Newborn screening | Spain (North West) | Newborns (N=14,600) | EA + G | FD: 1 in 7575 newborn males | |
| Hopkins et al 2018 ¹⁴ | Newborn screening | USA (Missouri) | Newborns (N=308,000) | EA + G | | FD: 1 in 3277 |
| Hult et al 2014 ¹⁵ | Retrospective | Sweden | N=23 cases* | EA | | FD: 1 in 90,000 births |
| Hwu et al 2009 ¹⁵ | Newborn screening | Taiwan | Consecutive newborns (N=171,977) | EA + G | <ul style="list-style-type: none"> Classic FD: ~1 in 22,570 newborn males FD: ~1 in 40,840 newborn females FD: ~1 in 1250 newborn males (86% of which carry cardiac variant IV54+919G-A) | |
| Inoue et al 2013 ¹⁶ | Newborn screening | Japan | Newborns (N=21,170) | EA + G | FD: 1 in 7057 newborns | |
| Navarrete et al 2015 ¹⁷ | Newborn screening | Mexico | Newborns (N=10,853) | NR | FD: 1 in 2713 newborns | |
| Navarrete-Martinez et al 2017 ¹⁸ | Newborn screening | Mexico | Newborns (N=20,018) | EA + G | FD: 1 in 4003 newborns; 1 in 2048 newborn males | |
| Lin et al 2009 ¹⁹ | Newborn screening | Taiwan-Chinese population | Newborns (N=110,027) | EA + G | <ul style="list-style-type: none"> FD: 1 in 1368 newborn males Cardiac variant IV54+919G-A: ~1 in 1600 newborn males | |
| Mechtler et al 2017 ²⁰ | Newborn screening | Austria | Consecutive newborns (N=34,736) | EA + G | Variants characteristic of FD: 1 per 3859 births | |
| Pasquini et al 2017 ²¹ | Newborn screening | Italy (Tuscany/Umbria) | Newborns (N=52,592) | EA + G | FD: 1 per 3757 newborns | |
| Pinto et al 2004 ²² | Retrospective | Portugal (Northern) | N=1 case* | EA + G | FD: ~1 per 833,000 live births | |
| Scott et al 2013 ²³ | Newborn screening | USA (Washington) | Newborn males (N=54,800) | EA + G | FD: 1 in 7800 newborn males | |
| Spada et al 2006 ²⁴ | Newborn screening | Italy (North West) | Consecutive newborn males (N=37,104) | EA + G | <ul style="list-style-type: none"> Late-onset FD: ~1 in 3100 newborn males Classic FD: ~1 in 37,000 newborn males | |
| Van der Tol et al 2014 ²⁵ | Systematic review | Various | Newborns (N=397,271) | Mostly EA | Pooled <i>GLA</i> variants: 1 in 2597 newborns* | |

*Extrapolated using the number of live births in the ascertainment period; †calculated but not reported in paper per se. α -Gal A, α -galactosidase A; EA, α -Gal A enzyme activity; FD, Fabry disease; G, genetic; NR, not reported.

Figure 2. Prevalence of Fabry disease-associated *GLA* gene and α -Gal A variants identified by newborn screening



- In Taiwanese and Chinese-Taiwanese populations, the incidence of a Fabry disease cardiac variant (IV54+919G-A) was ~1 in 1460 to ~1 in 1600 newborn males.^{15,19} However, further studies of the IV54 late-onset phenotype are needed to understand its natural history.
- Classic male Fabry disease prevalence in Italy was consistent with previous reports of 1 per 22,000 to 1 per 40,000 in the general male population,²⁶ reported as 1 per 22,570 newborn males and 1 per 37,000 newborn males in Taiwan and Italy, respectively.^{15,24}
- Lin and colleagues¹⁹ acknowledged that detecting hotspot mutations is preferred to lymphocyte enzyme activity to screen females because enzyme activity fails to predict disease severity in a particular organ.¹⁹
- In our literature search, a single newborn screening study reported the prevalence of Fabry disease-associated *GLA* variants among newborn females specifically, showing a significantly lower incidence than male newborns.¹⁵

LIMITATIONS

- This review was limited to studies published in English language. It is possible that relevant studies published in other languages were not retrieved.
- Unlike a systematic literature review, the present literature search was not subjected to a critical appraisal of the risk of bias. However, a targeted review approach can still be considered a rigorous and transparent method to synthesise evidence and provide insight into the incidence and prevalence of Fabry disease in newborn, paediatric and adolescent populations.

DISCUSSION

- Overall, Fabry disease epidemiology estimates are extremely heterogeneous between different countries and, while data are emerging for newborns based on gene and enzyme variants, epidemiological studies are particularly lacking in paediatric and adolescent populations.
- There is also a risk that estimates of *GLA* variant prevalence in females may be underestimated in screening programmes relying on α -Gal A enzyme assays.²⁷

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CONCLUSIONS

- Our findings support the importance of continued and wider implementation of newborn screening studies to better understand the epidemiology of Fabry disease-associated *GLA* variants
 - Care should be taken not to directly extrapolate the incidence and prevalence of *GLA* gene and α -Gal A variants to the clinical presentation of Fabry disease
 - Newborn screening results should therefore complement post-screening follow-up to ensure early symptoms are detected, enabling timely diagnosis and therapeutic intervention in the proportion who later present with symptomatic Fabry disease.
- Screening criteria for α -Gal A may need to be refined to ensure that females are not missed in newborn screening programmes.

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