

Of the importance of the clinical phenotypes in the interpretation of the studies dealing with Fabry disease

Wladimir Mauhin, Olivier Lidove, Olivier Benveniste

▶ To cite this version:

Wladimir Mauhin, Olivier Lidove, Olivier Benveniste. Of the importance of the clinical phenotypes in the interpretation of the studies dealing with Fabry disease. Orphanet Journal of Rare Diseases, 2019, 14 (1), pp.4. 10.1186/s13023-018-0979-z. hal-0.03523518

HAL Id: hal-03523518 https://hal.sorbonne-universite.fr/hal-03523518v1

Submitted on 12 Jan 2022

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

LETTER TO THE EDITOR

Open Access

Of the importance of the clinical phenotypes in the interpretation of the studies dealing with Fabry disease



Wladimir Mauhin^{1,2*}, Olivier Lidove^{1,2} and Olivier Benveniste²

Abstract

Fabry disease (OMIM #301500) is an X-linked disorder caused by alpha-galactosidase A deficiency with two major clinical phenotypes: classic and non-classic of different prognosis. From 2001, enzyme replacement therapies with agalsidase alfa and beta have been available. In this letter we underline the different clinical and technical considerations the readers have to be aware of to interpret the results of studies dealing with Fabry disease and anti-agalsidase antibodies. We reaffirm that antibodies preferentially develop in the severe classic Fabry phenotype, which can mislead into interpreting that antibodies are associated with much severe clinical events.

Keywords: Fabry disease, Antibodies, Inhibition, Phenotype

Dear Editors

We read with interest the letter from Lenders et al. concerning our recent article entitled "Deep characterization of the anti-drug antibodies developed in Fabry disease patients, a prospective analysis from the French multicenter cohort FFABRY" [1, 2]. In the letter, our main message seems to have been eluded: the development of anti-drug antibodies (ADAs) depends on the clinical phenotype (ADA-positivity in classic patients 58.6% vs 6.7% in non-classic patients, p < 0.001). Also, with the limits of a time-point study, after stratification on the clinical phenotype, we did not observe any obvious clinical event associated with the presence of ADAs. It is essential to remind that patients with a classical phenotype are more prone to develop a severe renal disease. In our cohort, all the kidney transplanted patients belonged to the classic group, independently from any ADA (see Fig. 1). Also we should have mentioned that 2 ADA-positive patients had benefited from a renal graft before the introduction of enzyme replacement therapy (ERT), suggesting an obvious severe disease prior to the development of antibodies. Concerning the exposure to ERT, the letter

We agree with Lenders and colleagues that purifying IgG subclasses could bring essential information concerning immunogenicity as a first approach. Also, it appears that this has not been performed in the referenced paper [3] where authors used purified total IgGs. We also agree with the authors that ADAs do not possess a mandatory neutralizing activity. This is the reason why we think that inhibition assays should only be performed after a first step using an immune-based assay such as an ELISA. Our goal was to study all ADAs, neutralizing and non-neutralizing. We may have to clarify that we did perform inhibition assay in all the men, contrary to what is mentioned in the letter. As expected, any of the antibody-negative serum was associated with enzymatic inhibition (Fig. 4a). It should also be reminded to readers that there is no consensus for inhibition assay and that the percentage of enzyme inhibition depends on the concentrations of

²Sorbonne Université, INSERM, UMR 974, Centre of Research in Myology, Association Institut de Myologie, Pitié-Salpêtrière University Hospital, 75013 Paris, France



mentions that "more ADA-positive patients were treated with agalsidase beta", which is wrong: as mentioned in our article, there was no difference in terms of seroprevalence in the different treatment group (alfa 30.8%, beta 44.4%, alfa and beta 42.9%, p = 0.7). There was also no difference in the mean infused dose received by patients during their whole exposure to ERT (ADA-positive vs ADA-negative patients 0.43 mg/kg vs 0.64 mg/kg, p = ns).

^{*} Correspondence: wladmauhin@gmail.com

¹Internal Medicine Department, Groupe hospitalier Diaconesses Croix Saint Simon, 125 rue d'Avron, 75020 Paris, France

ERT used in the protocol of the inhibition assay. Therefore, there is a need to standardize the protocol and the threshold retained to define inhibition.

In summary, we cannot conclude that antibodies (not only neutralizing) are associated with clinical events in our cohort in this time-point study.

Abbreviations

ADA: Anti-drug antibodies; ERT: Enzyme replacement therapy

Funding

Not applicable.

Availability of data and materials

The datasets generated and/or analysed during the current study are not publicly available due to the individual person's data that are involved but are available from the corresponding author on reasonable request.

Authors' contributions

WM, OL and OB designed the study, performed the experiments, interpreted the data, drafted the manuscript and approved this final version.

Ethics approval and consent to participate

Legal authorizations were obtained from the Comité consultatif sur le traitement de l'information en matière de recherche dans le domaine de la santé (n°14.324bis) and the Comité de Protection des personnes Paris VI, according to the relevant French legislation.

All patients signed written consent after specific oral and written information, for this research and its publication.

Consent for publication

All patients signed written consent after specific oral and written information, for this research and its publication.

Competing interests

Inserm U974 research team received financial support from Shire, AFM-Téléthon, SNFMI (Société Nationale Française de Médecine Interne) and VML (Vaincre les Maladies Lysosomales) for this study.

Wladimir Mauhin: Travel fees and accommodations: Shire, Orphan Europe, Amicus, Sanofi-Genzyme, honorarium: Shire.

Olivier Lidove: Travel fees, accommodations and honorarium: Shire, Sanofi-Genzyme; honorarium: Amicus.

Olivier Benveniste: Travel fees Shire, LFB, CSL Behring; honorarium: Novartis, Neovacs, LFB.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Received: 6 December 2018 Accepted: 12 December 2018 Published online: 07 January 2019

References

- 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 multicenter cohort FFABRY. Orphanet J Rare Dis 31 juill. 2018;13(1):127.
- Lenders M, Schmitz B, Brand S-M, Brand E. Neutralizing anti-drug antibodies in Fabry disease have no obvious clinical impact? Orphanet J Rare Dis. 2018; 13(1):171.
- Lenders M, Schmitz B, Brand S-M, Foell D, Brand E. Characterization of drugneutralizing antibodies in patients with Fabry disease during infusion. J Allergy Clin Immunol. 2018;141(6):2289–2292.e7.

Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

