Optimal therapy in Gaucher disease

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Abstract: Gaucher disease (GD), the inherited deficiency of the lysosomal enzyme glucocerebrosidase, presents with a wide range of symptoms of varying severity, and primarily affects the skeletal, hematologic and nervous systems. To date, the standard of care has included enzyme replacement therapy with imiglucerase. Although imiglucerase is highly effective in reversing the visceral and hematologic manifestations, skeletal disease is slow to respond, pulmonary involvement is relatively resistant, and the CNS involvement is not impacted. Because of the recent manufacturing and processing problems, the research and development of alternative therapeutics has become more pressing. The divergent phenotypes and the heterogeneity involving different organ systems implicates the involvement of several pathological processes that include enzyme deficiency, substrate accumulation, protein misfolding, and macrophage activation, that differ in each patient with GD. Thus, the therapy should be tailored individually in order to target multiple pathways that interplay in GD.

Keywords: glucocerebrosidase, enzyme replacement therapy, substrate reduction therapy, protein misfolding and chaperone therapy, macrophage

Introduction
Gaucher disease (GD), the inherited deficiency of the lysosomal enzyme glucocerebrosidase ([GC] enzyme commission number EC 3.2.1.45), presents with a wide range of symptoms of varying severity, and primarily affects the skeletal, hematologic and nervous systems. Clinically, there are three types of GD based upon the presence and progression of the neurological symptoms. Absence of central nervous system (CNS) involvement is considered mandatory for a diagnosis of the adult form or type 1 GD, also known as nonneuronopathic GD. When there is neuronopathic involvement, the distinction between infantile, type 2 or juvenile and the type 3 form of GD, is made in the basis of age of onset and progression of CNS symptoms. While an intermediate phenotype between types 2 and 3 is also described.1

The gene encoding GC (GBA, MIM# 606463) and its homologous pseudogene are located on chromosome 1q21.2 To date, more than 250 different GBA mutations, including missense, nonsense, insertions or deletions that lead to frameshift or in-frame alterations, and splice junction mutations have been identified. Recombination events with a highly homologous pseudogene downstream of the GBA locus have also been reported.3 Although certain mutations, especially N370S are associated only with adult or nonneuronopathic forms, the validity of a phenotype prediction based upon genotype is quite limited. Likewise, although the homozygosity for L444P mutation...
results in neuronopathic GD, a milder clinical phenotype could be observed with very low residual enzymatic activities. Remarkably, phenotype and disease severity may vary among siblings, even in identical twins, and among patients carrying same mutations, a wide range of phenotypes and residual enzymatic activity are observed.

GC is a membrane-associated lysosomal glycoprotein. After synthesis and initial folding at a neutral pH in the endoplasmic reticulum, it is targeted to the lysosomes, where the acidic pH provides the optimal environment for stable folding and function. The majority of GBA mutations lead to the synthesis of GC with decreased catalytic function and/or stability. While it is assumed that disease severity depends on residual GC activity, this has been difficult to prove for most GBA mutations.

GC deficiency results in the accumulation of the lipid substrate glucosylceramide. Glucosylceramide is the common intermediate resulting from lysosomal degradation of gangliosides and globo- sides, primarily in cells derived from monocyte-macrophage origin. The hallmark “Gaucher cells” are glucosylceramide-laden macrophages, observed in many organs, although they are predominant in bone marrow, liver, spleen, and the parenchyma of lymph nodes. The accumulation of glucosylceramide causes enlargement of liver and spleen. Splenomegaly is associated with hypersplenism leading to anemia and thrombocytopenia. Pancytopenia is also caused by replacement of the bone marrow. Other skeletal manifestations include decreased mineralization, pathological fractures, bone pain, avascular necrosis, and painful “bone crises”. There is primarily radiological evidence of interstitial lung involvement, and on rare occasions pulmonary hypertension can occur. Additionally in children growth failure is also a common feature.

This review will focus on the mechanisms interplaying in GD, and discuss current therapy and others, that are on the immediate horizon, that target different pathological pathways.

**Gaucher disease, the deficiency of the lysosomal enzyme glucocerebrosidase: recombinant enzyme therapy**

Human GC is encoded by a single locus (GBA) at chromosome 1q21. This lysosome-associated glycoprotein has 497 amino acids, and N-glycosylation at four or five different sites that is essential for its catalytic activity and half life. In the lysosome, GC catalyzes a two-step reaction that requires glycosylation of its active site by the substrate, followed by the release of beta-glucose by deglucosylation. To enhance *in vitro* activity, negatively charged phospholipids, and a naturally occurring 80 amino acid protein, saposin C are required. While there is evidence that GC physically interacts with saposin C, the exact mechanism of saposin C-mediated GC activation is not known.

The resolution of the X-ray crystal structure of human GC, followed by the demonstration of a nonreversible inhibitor, conduritol β-epoxide (CBE) bound GC structure helped to understand the effects of the tertiary structure on catalytic function, and spatial relationship of GBA mutations with the active site. The 3D structure with the CBE had confirmed E340 as the nucleophile and E235 as the acid-base catalyst. Although few mutations that are associated with a severe phenotype occur near the catalytic site, the majority of GBA variants are spread through the domains of the protein. However, there is no clear relationship between the position of GBA variants in the 3D structure and the disease severity. While the crystal structure is not available for GBA mutants; among the common mutations L444, which is mutated to proline or arginine, is located at the hydrophobic core of the protein, and is predicted to produce unstable protein. Whereas the mutation N370S has been suggested to flank the active site, and alter interaction with saposin C and anionic phospholipid membranes.

In 1964, De Duve introduced the concept of replacing the deficiency of a lysosomal enzyme. Proof of the general principles of enzyme replacement therapy (ERT) was developed in a model system using cocultivation of fibroblasts from patients with different types of mucopolysaccharidosis, that corrected the enzymatic deficiency. Early attempts to correct the enzymatic deficiency in 1970’s revealed the critical importance of N-linked oligosaccharide chain stability of the lysosomal enzymes, and the importance of cell-type specific uptake in different diseases. The recognition signal was initially identified as mannos, and *in vivo* clearance of the lysosomal enzymes from plasma was found to be mediated by mannos receptors of the reticuloendothelial system. ERT quickly advanced in GD, where macrophages were regarded as the primary source for the stored abnormal lipid. The first enzyme preparation purified from placental tissue did not yield any significant clinical response, although the total amount of stored lipid was shown to be reduced in liver samples. The placental enzyme that contained complex N-linked glycans was largely taken up by the asialoglycoprotein receptors present on hepatocytes. Further modification of N-linked oligosaccharide terminals to remove sialic acid residues, that exposed terminal mannos residues, was essential for the
internalization of the enzyme by the Gaucher macrophages. This modified high-mannose form of enzyme was readily recognized by mannose receptors abundant on the macrophages. The first placental derived enzyme product alglucerase (Cerezyme®; Genzyme Inc, Cambridge, MA, USA) was developed based on this principle, and was tested successfully on 12 patients with type 1 GD with improvements in hematological parameters, and the reduction in organ volumes. This was followed by two trials on 25 patients with imiglucerase (Cerezyme®; Genzyme Inc), a recombinant form of human GC, which differs from native GC with a substitution of histidine instead of arginine at position 495. Until the clinical use of ERT in 1990’s, the treatment of GD consisted only of palliative measures such as splenectomy and hip replacement. The development of ERT for GD is considered to be a triumph of translational medicine. ERT is a safe and well tolerated treatment modality with only rare side effects. Currently, there are more than 5000 patients with GD that are successfully being treated with imiglucerase.

In vivo, recombinant GC shows high-affinity targeting to the macrophages. In humans and mice, imiglucerase has been shown to exhibit similar postinjection distribution patterns. The uptake of the human enzyme is primarily by the reticuloendothelial cells of the liver and spleen. In mice, post-injection enzymatic activity peaks at about 20 minutes in the liver and spleen, followed by a biphasic decrease, with a half-life around 40–60 minutes and 12–14 hours respectively. While in bone marrow, the maximal enzymatic activity is measured at 40–60 minutes. The uptake and tissue distribution of radio-labeled recombinant enzyme after a bolus injection was followed in patients with type 1 GD. About half of the radio-labeled enzyme was rapidly cleared from blood with a half-life less than five minutes. 30% of the injected dose was taken up immediately by the liver, about 15% by the spleen, and the bone marrow, where the half-life was over 14 hours. The remainder was cleared slowly with a half-life of between 34–42 hours. When the enzyme was administered at a higher dose, the receptor-mediated uptake was saturated, and the clearance half-life of tracer enzyme was prolonged.

During the original clinical trial, imiglucerase was administered at a dose of 60 IU/kg every two weeks. There is a dose-dependent improvement in hematological and visceral parameters with alglucerase/imiglucerase. Patients with type 1 GD treated with doses more than 48 IU/kg, every two weeks, were suggested to have better clinical responses. However, lower dose higher frequency or higher doses administered at less frequent intervals could provide similar clinical efficacy. The therapeutic threshold for macrophage-targeted GC appears to be 10–15 IU/kg every 2 weeks. Although high dose therapy provides a faster clinical response, and should be considered for patients with a more aggressive disease, the initiation of ERT at a lower dose has also led to an improvement in all nonskeletal manifestations of GD. The effects of initiating ERT at a high dose, followed by progressive dose reductions, were compared with starting the therapy at a low dose only. In patients who were started on high-dose ERT, hemoglobin, acid phosphatase, and organ volume improved or remained unchanged at the end of each dose reduction, although the platelet count decreased when the dose of enzyme was reduced to 15 IU/kg. While in the treatment group that started at the minimal dose, the manifestations were similarly well controlled. In another study, there were no significant differences in clinical outcome when the recombinant enzyme was given at different dosing schedules including: 15 IU/kg once fortnightly or 2.5 IU/kg thrice weekly. Thus, smaller total doses of recombinant GC, administered frequently may have significant financial benefits, and could be considered in patients with stable type 1 GD. Similarly, even lower doses (7.5 IU/kg every two weeks) established good visceral and hematological response in adult patients with mild to moderate skeletal involvement. Despite some rare reports of treatment failures in patients with a higher disease burden, ERT administered every four weeks, was shown to be effective in most adult type 1 GD patients with stable and minimal disease. Currently, the recommendations for commencing imiglucerase treatment and subsequent dose adjustments are based on regular assessment and monitoring. Then, the maintenance of defined therapeutic goals, that include hematopoietic reconstitution, the reduction of liver and spleen volumes, together with the stabilization or improvement in skeletal findings is achieved on an individualized regimen. While tailoring therapy for each individual patient, several other parameters including the cost, and impact to the life style should also be taken into consideration.

In individuals with nonneuronopathic GD presentation during childhood years suggests a more severe phenotype. However many patients who are diagnosed in large-scale screenings, are mildly affected and may not require treatment. In a retrospective study, 56 children with nonneuronopathic GD, were followed up to 9 years, and only massive splenomegaly and growth retardation were found to be indications for ERT. In symptomatic children, however, early intervention aims to prevent the development of complications, and ensure normal growth and development through adolescence. In a large international cohort of children with GD type 1 on ERT, the longitudinal data supports the normalization of most clinical parameters including mean height, hemoglobin values...
and mean bone mineral density z scores. Several consensus statements for treatment and monitoring of children with GD, include regular assessments, and individualized dose adjustments based on clinical response. Although a lower maintenance dose of 30–35 IU/kg/month may be effective in most pediatric patients, more severely affected ones, especially L444P/L444P homozygotes, may require a higher maintenance dosage. In addition to clinical parameters, the follow up in this age group should also include evaluation of psychosocial, functional status and quality-of-life assessments.

Although imiglucerase is highly effective in reversing the visceral and hematologic manifestations, skeletal disease is slow to respond, pulmonary involvement is relatively resistant to the enzyme, and the CNS involvement is not impacted by ERT. Some patients with type 1 GD even may experience ongoing bone disease, despite increasing doses of ERT. In such patients the ERT dose should be adjusted in order to maintain adequate visceral control, although alternative strategies including the addition of bisphosphonates could be explored. There is great heterogeneity in response to ERT in patients with GD and symptomatic lung involvement. Clinically, some patients benefit significantly from enzyme therapy. However, pulmonary function or radiological findings do not usually normalize, despite the correction of hematological parameters and the reduction in organ volumes. Although worsening pulmonary pressures have been reported after the introduction of ERT, pulmonary hypertension is associated either with poor compliance, or the lack of ERT. An emerging concern has been whether ERT could prevent the progression of the neurological disease in neuronopathic GD. The guidelines for the management of the neuronopathic forms of GD were recently revised. In particular, the role of high-dose ERT (120 IU/kg of body weight every 2 weeks) was analyzed. In patients receiving high-dose ERT, there was a consistent progression of brain stem findings, concomitant with the deterioration of auditory brain stem responses. Although high dose ERT may be required to control severe systemic symptoms, it does not stabilize neurological symptoms, and thus, is not recommended in neuronopathic GD.

Other recombinant glucocerebrosidases, recently approved by the FDA or pending approval, include velaglucerase alpha (Shire HGT, Cambridge, MA, USA) and taliglucerase alpha (Protalix Biotherapeutics, Carmiel, Israel; Pfizer, NY, USA). Both have successfully completed phase III trials, and proven to be safe and clinically efficient. Although considered as biosimilars, the new recombinant GCs are not generic proteins, they are produced as novel biologicals. Imiglucerase is expressed in Chinese hamster ovary cells, where as taliglucerase is produced in transgenic carrot cells. In contrast to imiglucerase and taliglucerase, velaglucerase alpha, the gene-activated human GC, contains the native human enzyme sequence. On X-ray crystallography, all three glucocerebrosidases consist of three domains, and the active site is located in domain III, and E340 and E235 are the catalytic residues. Taliglucerase alpha demonstrates a high degree of homology to imiglucerase in its high-resolution three-dimensional structure, and has a bioactivity similar to imiglucerase. In clinical trials, velaglucerase displayed pharmacokinetic properties suggesting that the tissue distribution is linear to the doses administered. In patients, the serum concentration increased linearly with increasing doses from 15 IU/kg to 60 IU/kg, and the maximum serum concentration was recorded at the end of the first hour of the infusion. Although velaglucerase has a specific activity comparable with imiglucerase, the cellular uptake is suggested to be more efficient due to differences in glycosylation. Imiglucerase contains a chitobiose tri-mannosyl core glycan with fucosylation, while velaglucerase predominantly displays high-mannose type glycans with nine mannose units. The differences in glycosylation are suggested to affect better internalization into human macrophages. Taliglucerase is targeted to the storage vacuoles using a plant-specific C-terminal sorting signal, that naturally contains terminal mannose residues in its complex glycans. Thus, taliglucerase does not require exposure of mannose residues in vitro, which is a requirement for the production of other recombinant GCs. Nevertheless, higher number of mannose residues may result in increased immunogenicity due to the presentation of nonmammalian glycans. However, there have been no clinical adverse effects, nor increased frequency of neutralizing antibodies associated with taliglucerase during safety and pharmacokinetics studies. Recently, there has been serious problems in the manufacturing and processing of imiglucerase, and availability of alternative therapeutics has become more pressing than ever. Having different enzymes for therapeutic use will provide an opportunity to re-evaluate indications and initiation of ERT, management of nonresponders, and a cost–benefit maintenance therapy.

Gaucher disease, a storage disorder of the lysosomes: substrate reduction therapy (SRT)

The pathophysiology of lysosomal storage diseases is not completely understood and there is no doubt that the abnormal substrate accumulation, the triggering
secondary biochemical and cellular events, is one of the primary the pathogenic factors. In GD, glucosylceramide accumulates in the lysosomes, it is not known whether the cellular pathology is related to the lysosomal dysfunction secondary to storage, or whether glucosylceramide escapes from the lysosome and interacts with other organelles. In macrophages, for example, glucosylceramide accumulation is suggested to cause enhanced phagocytic activity.55 Glucosylceramide accumulates in tissues, and is elevated in the plasma of patients with GD, the amount is not significant.56 While ceramide accumulates in tissues, and is elevated in the plasma of patients with GD type 3, miglustat unfortunately, does not decrease in chitotriosidase levels have been observed in patients with GD type 1, who are unable or unwilling to be on ERT.57 However, significant side effects such as loose stools, tremor and peripheral neuropathy have limited its therapeutic use.58 Currently attempts are ongoing to develop other glucosylceramide inhibitors, and a phase III clinical trial is being conducted to evaluate the efficacy, safety, and pharmacokinetics of a novel SRT molecule in adult patients with GD type 1 (ClinicalTrials.gov identifier: NCT00891202). Unlike recombinant enzymes, SRT agents are small molecules that can cross the blood-brain barrier.59 Although improved pulmonary function, and a significant decrease in chitotriosidase levels have been observed in patients with GD type 3, miglustat unfortunately, does not appear to have any significant benefits on the neurological manifestations of GD.60

Glycosphingolipids affect a variety of cellular functions, and long term SRT may theoretically alter several pathways in the cell, and therefore, should be approached with great clinical care.61

**Gaucher disease, a protein misfolding disorder: pharmacological chaperones and enzyme enhancement therapy (EET)**

The majority of GBA mutations are missense, leading to the synthesis of GC with decreased catalytic function and/or stability.3 At the cellular level, missense mutations can result in altered processing or a conformational change in the mutant protein leading to a toxic gain-of-function.52,63 Protein misfolding contributes to the pathophysiology of several genetic and sporadic diseases. A prime example for proteinopathies is the adult-onset neurodegenerative diseases, where diverse molecular mechanisms result in the abnormal accumulation of proteins in the nervous system. Parkinson disease is the prototype adult-onset neurodegenerative disorder, characterized by the abnormal accumulation alpha-synuclein that results in intraneuronal aggregates and cell death.64 Clinical, genetic and neuropathologic observations have shown that GBA mutations constitute a risk factor for the development of parkinsonian manifestations, thus leading to the hypothesis that protein misfolding may play a role in the phenotypes associated with GD.65-67

“Pharmacological chaperones” are small molecules that bind to misfolded proteins, and assist their correct refolding and/or maturation. Chaperone or EET is an attractive approach because of its potential for simple oral administration, penetration of the blood–brain barrier, and its low cost. EET has been proposed for a number of lysosomal storage disorders, including GD, Sandhoff, Fabry, and Tay-Sachs diseases.68,69 The mechanism of action for these compounds is competitive binding to the active site of the mutant enzyme, facilitating proper folding and trafficking to the lysosome, where endogenous substrate displaces the chaperone and enzyme activity is restored.70 Most chaperones studied to date are enzyme inhibitors in the structural class of imino sugars or similar analogs of the natural substrate, glucosylceramide.71 Imino sugars have been shown to increase the cellular activity of the N370S mutant form of GC, as well as the wild-type enzyme.72 In cell based systems, where fibroblasts from patients are cultured with the chemical chaperone N-nonyl-deoxynojirimycin, the enzymatic activity of the GC variant, N370S, is increased. Similarly G202R, a GC variant retained in the endoplasmic reticulum, is also shown to be amenable to chemical chaperoning.73 However, imino sugar derivatives are nonspecific and have relatively short half-lives in cells, and these molecules can inhibit other glycolipid and glycoprotein processing enzymes.74 Several studies have demonstrated that isofagomine (IFG), an imino sugar that binds to the active site, enhanced the folding, transport, and stability of several GC variants, including L444P.75 IFG was found to have little or no inhibitory activity against other glucosidases, and further treatment of wild type fibroblasts with high doses did not alter the processing of newly synthesized N-linked oligosaccharides.76 These findings supported the further evaluation of IFG as a potential therapeutic
agent in GD. Despite promising in vivo results, and safety data (Clinical Trials.gov Identifier: NCT00433147), clinically meaningful improvements in key measures of the disease were not adequate. There are no plans to advance this chaperone further into a phase III trial. In addition to imino sugars, calcium channel blockers are suggested to act as pharmacological chaperones through altered calcium homeostasis, and are shown to partially restore mutant GC activity. While diltiazem exhibited the properties of a pharmacological chaperone at neutral pH, of the endoplasmic reticulum in cell-based systems, in animal studies these effects were minor.

Screening large compound libraries by quantitative high-throughput screening (qHTS) provides comprehensive information on the potency, efficacy, and structure-activity relationships of active compounds, facilitating identification of leads for medicinal chemistry optimization. qHTS was used to rapidly identify different nonsugar GC inhibitors. Similarly, screening a library of FDA-approved drugs using a thermal denaturation assay identified Ambroxol (a drug used to treat airway mucus hypersecretion and hyaline membrane disease in newborns) as a pH-dependent, mixed-type inhibitor of GC. Ambroxol demonstrates the characteristics of an EET agent through its interaction with both active and nonactive sites, and may have the potential to move towards clinical trials.

Although EET is promising to treat CNS manifestations, in vitro studies have shown that pharmacological chaperones are mostly mutation specific, and they may not be suitable for every patient with GD. Long-term clinical trials are required to test the in vivo effects of different chaperones on the specific disease pathology that could be associated with protein misfolding.

**Gaucher disease, a macrophage disorder**

GD was described more than a century ago by the identification of lipid engorged macrophages, referred to as “Gaucher cells”, were periodic acid Schiff-positive cells with wrinkled-tissue paper like appearance. GD has been regarded primarily as a macrophage disorder; however, current knowledge renders this classification too simplistic. Gaucher cells are not inert storage cells, they are metabolically active and secrete various proteins. In the plasma of patients with GD, the levels of macrophage/monocyte activation markers such as soluble CD14 and CD 163 are elevated. Gaucher cells strongly express markers of alternative activation, such as interleukin-1 antagonist and CCL18, which are anti-inflammatory molecules. Other biochemical correlates of GD represent hydrolases, localized to storage cells and/or surrounding macrophages, such as tartrate-resistant acid phosphatase and angiotensin-converting enzyme. These biomarkers, however, are nonspecific, and are altered in a number of disorders that affect the reticuloendothelial system. Furthermore, the overlap between the levels of hydrolases between patients with GD and healthy controls restrict their clinical use.

In the plasma of symptomatic GD patients, biochemical investigations demonstrated a more than 1000-fold increase of a class of chitinases, named “chitotriosidase”. Chitotriosidase originates from macrophages, and stored lipid is the major inducer. Macrophages are also the primary source for another cytokine, CCL18, which has been found to be elevated in GD. Both these surrogate markers are not organ specific, although they may reflect overall disease burden, and respond to therapeutic response. While the exact contribution of chitotriosidase and CCL18 to the disease pathophysiology remains to be elucidated, the pulmonary, skeletal and immune system involvement in GD may have a closer association with the activation and release of different cytokines from the macrophage-monocyte system. During the chronic inflammatory process, as is observed in some patients with GD, the balance between bone formation and resorption is skewed toward osteoclast-mediated bone resorption, which may explain osteopenia commonly observed in patients with GD. Macrophage activation is the common denominator in multiple disorders, characterized by radiological evidence of interstitial pulmonary involvement. While chitotriosidase is one of the established biomarkers for GD, it has been recently used as an indicator in alveolar lung diseases, such as sarcoidosis and miliary tuberculosis. Chitotriosidase levels are shown to be significantly higher in patients with idiopathic interstitial lung disease.

Surrogate markers in GD serve only as indicators of macrophage activation, and should not be used as diagnostic tools. Macrophages participate in the autoregulatory loop of the inflammatory process, and produce a wide range of biologically active molecules participating in both the beneficial and detrimental outcomes in inflammation. The inflammatory component in GD may play a role in the development and progression of bone, pulmonary and neurological complications. Therapeutic interventions that target macrophages and their products may open new avenues in the treatment of GD.

**Conclusion**

Although GD is considered to be a “simple Mendelian disorder”, divergent phenotypes and the heterogeneity involvement of different organ systems in each patient with GD implicate the role
of different pathological pathways. Thus the optimal therapeutic expectations may differ in each patient with GD. Despite better tissue penetration and alternative mechanisms of action of novel therapeutics what is potentially achievable and what is likely to remain irreversible in GD manifestations will be contingent upon multiple factors such as genetic and environmental modifiers. The bone and secondary neurological manifestations in adult-onset GD may serve as examples for some perceived limitations of clinical management in GD. Although the ultimate aim is to treat all skeletal manifestations, such as the reversal of osteopoenia and to prevent and treat bone crises, a clinician who follows a number of patients with GD will be aware that some patients will progressively lose bone mineral density or develop new bone lesions, despite a maximum dose ERT, diet supplements or weight-bearing exercises. The establishment of the association between GBA mutations and the development of parkinsonian manifestations have brought new challenges for both the clinician and researchers. There are several lines of evidence that suggest there is an association between GBA mutations and development of Parkinson disease pathology.60,65 Although the risk of Parkinson disease is suggested to be increased, especially among individuals carrying “severe” GBA mutations, the penetrance and again the role of modifiers are not known for each GBA alteration.66 Furthermore, 99% of individuals carrying GBA mutations either in hetero- or homozygote states will not develop parkinsonian manifestations, limiting the genetic counseling even in “at risk” individuals in families with more than one person affected. The current recommended therapy for GBA-associated parkinsonism is limited to the standard parkinsonian therapy, when the symptoms are L-dopa responsive, and testing for GBA mutations remains an academic endeavor more than that of a risk and prognostic determinant.

The future of the treatment of GD will include an individualized approach, employing a combination of therapies that can address the effects of enzyme deficiency, storage of abnormal lipid substrate resulting in lysosomal dysfunction, protein misfolding and the inflammatory processes.

Disclosure
The author reports no conflicts of interest relevant to this research.

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