Skeletal muscle effects of antisense oligonucleotides targeting glycogen synthase 1 in a mouse

model of Pompe disease

Lan Weiss[1*], Michele Carrer[2*], Alyaa Shmara[1], Cheng Cheng[1], Hong Yin[3], Lac Ta[1], Victoria

Boock[1], Yasamin Fazeli[1], Mindy Chang[1], Marvin Paguio[1], Jonathan Lee[1], Howard Yu[1], Angela

Martin[1], Nina Raben[4], John Weiss[3], Tamar Grossman[2†], Paymaan Jafar-nejad[2], Virginia Kimonis[1#]

Affiliations:

[1] Division of Genetics and Genomic Medicine, Department of Pediatrics, University of California,

Irvine, CA 92868, USA

[2] Ionis Pharmaceuticals, Inc., Carlsbad, CA, 92010, USA

[3] Department of Neurology, University of California, Irvine, CA, 92868, USA

[4] Cell and Developmental Biology Center, National Heart, Lung, and Blood Institute, National

Institutes of Health, Bethesda, MD, 20892, USA. Current address: M6P Therapeutics, St. Louis,

MO

[†] Current address: The Janssen Pharmaceutical Companies of Johnson & Johnson, San Diego, CA,

92121, USA

* These authors contributed equally to this work

[#] To whom correspondence should be addressed:

Virginia Kimonis, MD

Division of Genetics and Genomic Medicine

Department of Pediatrics, Neurology and Pathology

University of California, Irvine

Tel: (949) 824-0571

Fax: (949) 824-0171

Email: vkimonis@uci.edu

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rhGAA: recombinant human GAA; SRT: substrate reduction therapy; ERT: enzyme replacement

therapy; PD: Pompe disease

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ABSTRACT

Pompe disease (PD) is a progressive myopathy caused by the aberrant accumulation of glycogen in skeletal and cardiac muscle resulting from the deficiency of the enzyme acid alpha-glucosidase (GAA). Administration of recombinant human GAA as enzyme replacement therapy (ERT) works well in alleviating the cardiac manifestations of PD but loses sustained benefit in ameliorating the skeletal muscle pathology. The limited efficacy of ERT in skeletal muscle is partially attributable to its inability to curb the accumulation of new glycogen produced by the muscle enzyme glycogen synthase 1 (GYS1). Substrate reduction therapies aimed at knocking down GYS1 expression represent a promising avenue to improve Pompe myopathy. However, finding specific inhibitors for GYS1 is challenging given the presence of the highly homologous GYS2 in the liver. Antisense oligonucleotides (ASOs) are chemically modified oligomers that hybridize to their complementary target RNA to induce their degradation with exquisite specificity. In the present study, we show that ASO-mediated Gys1 knockdown in the $Gaa^{-/-}$ mouse model of PD led to a robust reduction in glycogen accumulation in skeletal and cardiac muscle. In addition, combining Gys1 ASO with ERT further reduced glycogen content in muscle, eliminated autophagic buildup and lysosomal dysfunction, and improved motor function in $Gaa^{-/-}$ mice. Our results provide a strong foundation for further validation of the use of Gys1 ASO, alone or in combination with ERT, as a therapy for PD. We propose that early administration of Gys1 ASO in combination with ERT may be the key to preventative treatment options in PD.

INTRODUCTION

Glycogen storage disease type II, also called Pompe disease (PD) is an autosomal recessive disorder affecting between 1 in 17,000 and 1 in 40,000 individuals[1]. PD is caused by the excessive accumulation of glycogen in tissues resulting from inactivating mutations in the GAA gene, which encodes acid alpha-glucosidase, an enzyme responsible for the lysosomal degradation of glycogen. The pathological accumulation of glycogen affects many tissues, but it is by far most impactful in the heart and skeletal muscle of Pompe patients. Indeed, the lack of GAA enzyme activity, or extremely low levels (1–2% of normal), are associated with a fatal infantile cardiomyopathy, profound weakness, and skeletal muscle pathology, whereas 10-20% of normal GAA levels lead to skeletal myopathy with childhood or adult onset[2,3]. To understand the role of GAA in Pompe disease etiology, a mouse model has been generated by targeted deletion of the murine Gaa gene. The Gaa knockout (Gaa$^{-/-}$) mouse recapitulates Pompe disease manifestations, including the accumulation of glycogen in skeletal and cardiac muscle, the autophagy defects, and the consequent myopathy[4].

The standard of care for Pompe patients is enzyme replacement therapy (ERT) using recombinant human (rh) GAA (Lumizyme®, or Myozyme®, Sanofi Aventis US LCC), which became commercially available in 2006. ERT has shown positive results in improving the myopathy and cardiomyopathy associated with PD, and therefore has been used to help stabilize or partially improve patient outcomes[5-7]. However, ERT has limited efficacy in the skeletal muscle, as many patients continue to develop progressive muscle weakness despite receiving rhGAA enzyme[8-10]. Multiple factors contribute to the poor efficacy of ERT in the skeletal muscle of Pompe patients, particularly when significant disease pathology is present. Some of the limitations affecting ERT are associated with the complexity of delivering a relatively large protein to the muscle tissue.

Indeed, when administered via intravenous infusion, the recombinant GAA enzyme circulates to

the liver, which acts as a major "sink" and precludes more than 80% of the administered protein

from reaching the skeletal muscle[11]. Furthermore, ERT relies on the cation-independent mannose-

6-phosphate receptor (CI-MPR), which is expressed at low levels on the membrane of skeletal

muscle cells[7]. In addition, dysregulation of autophagy, and the consequent impairment of the

autophagosome-lysosome fusion in Pompe myofibers, further contribute to the reduced efficacy

of ERT. Ultimately, the failure of autophagy leads to inefficient trafficking of the therapeutic

enzyme to the lysosome, limiting its ability to clear the glycogen[12]. This phenomenon is especially

evident in type II muscle fibers, which are particularly refractory to ERT[10]. Previous studies

performed in mice found upregulated protein levels of enzymes involved in glucose uptake and

cytoplasmic glycogen synthesis in skeletal muscle from mice with Pompe disease, including

glycogenin (GYG1), glycogen synthase (GYS1), glucose transporter 4 (GLUT4), glycogen

branching enzyme 1 (GBE1), and UDP-glucose pyrophosphorylase (UGP2) suggesting a positive

feedforward loop for cellular glycogen accumulation[13]. Administration of rhGAA to the Pompe

mice normalized not only muscle glycogen levels but also G6P, hexokinase, GS, and glycogenin

levels[14]. Thus, it would appear that correction of one aspect of the glycogen metabolic pathway

(lysosomal glycogen) results in the correction of multiple other components. Studies in patient

biopsies however found that extralysosomal glycogen accumulation in muscle biopsies from late-

onset patients was not cleared by ERT, indicating the importance of an acidic pH environment[15].

Glycogen synthase (GYS) enzymes catalyze the addition of glucose residues to the growing

glycogen molecules by formation of α-1,4-glucosidic linkages. Two paralogous GYS genes, GYS1

and GYS2, exist in mice and humans. GYS2 expression is restricted to the liver, where it plays an

essential role in the maintenance of proper body glucose homeostasis[16]. GYS1 has a broader

expression profile, but it is particularly abundant in skeletal muscle and heart[16]. Decreasing GYS1

expression represents a potential therapeutic approach for PD, based on the principle of substrate

reduction. Indeed, limiting the synthesis of new glycogen would result in attenuated glycogen

accumulation in the muscle tissue. Genetic ablation of murine Gys1 provided proof of principle

validation for substrate reduction therapy (SRT) in vivo. Knocking down Gys1 led to restoration

of muscle functionality in the Gaa$^{-/-}$ mouse model[17]. Similarly, intramuscular injection of

recombinant adeno-associated virus-1 (AAV-1) vector expressing short hairpin ribonucleic acid

(shRNA) targeted to Gys1 has been shown to reduce glycogen accumulation in the gastrocnemius

of newborn Gaa$^{-/-}$ mice [18]. Altogether the preclinical evidence suggests that reducing GYS1 can be

a potential therapeutic approach in Pompe disease, either alone or in combination with ERT.

However, GYS1 and GYS2 show high degree of homology, which poses a complex challenge in

the identification of safe, isoform-specific inhibitors[19]. It is important to note that no approved

small molecule inhibitor is clinically available for the treatment of Pompe disease.

Antisense oligonucleotides (ASOs) are short synthetic nucleic acids that hybridize to

complementary target RNAs in a sequence-specific manner using Watson-Crick base pairing. By

doing so, ASOs can influence RNA processing and regulate protein expression via multiple

mechanisms, including RNase H1-mediated mRNA degradation, and splicing modulation[20-24].

Over the past two decades ASOs have emerged as both powerful research tools and therapeutic

molecules[25-29]. The intrinsic properties of ASOs make them the ideal pharmacological tool to

reduce muscle GYS1 expression with exquisite specificity.

Here, we show that systemic treatment with a phosphorothioate (PS) 2'-constrained ethyl (cEt)-

modified gapmer ASO targeting murine Gys1 not only reversed the excessive glycogen

accumulation, but also corrected the autophagic defects present in the skeletal muscle of Gaa$^{-/-}$

mice. The ASO-mediated reduction of GYS1 protein, especially when combined with ERT, led to

restoration of muscle function in Pompe mice, thus underscoring the in vivo potential of our

approach.

RESULTS

Progressive glycogen accumulation and muscle weakness in the Gaa$^{-/-}$ mouse model

We utilized the Gaa knockout mouse model to study the effects of lowering Gys1 expression on

Pompe myopathy in vivo. The Gaa$^{-/-}$ mouse was generated and studied by Raben et al. (1998)[4]. To

inform the design of appropriate prevention and reversal treatment paradigms, we characterized

the progressive accumulation of glycogen in the skeletal and cardiac muscle of the Gaa$^{-/-}$ mice

from our colony. We utilized two methods to measure glycogen content in the tissues: a

colorimetric biochemical assay, and periodic acid-Schiff (PAS) histochemical staining[4]. As

expected, we observed higher amounts of glycogen in the skeletal muscle of Gaa$^{-/-}$ mice compared

to wild type mice (Figure 1A). Remarkably, the glycogen levels in quadriceps muscle were already

elevated in one-month-old Gaa$^{-/-}$ mice, suggesting that the mutant mice start to accumulate

glycogen in skeletal muscle at an early age.

The accumulation of glycogen increased progressively over time, from 0.26, to 0.41, and 0.55 μg

glycogen per μg of total protein in quadriceps muscle homogenate from one-, two-, and three-

month-old Gaa$^{-/-}$ mice, respectively (Figure 1A). Similar to previous reports [4,30,31], the excessive

glycogen accumulation in the skeletal muscle was associated with a progressive decline in motor

function in aging Gaa$^{-/-}$ mice, as assessed by the rotarod test (Figure 1B).

Different levels of ASO-mediated Gys1 knockdown lead to the proportional amelioration of

aberrant glycogen accumulation in the skeletal muscle of young Gaa$^{-/-}$ mice

Through in vitro and in vivo screening, we identified two ASOs targeting mouse Gys1 that were potent and well tolerated in mice (Supplementary Figure 1, 2). When dosed systemically to wild type mice via subcutaneous injection for six weeks, Gys1 ASO#1 and ASO#2 resulted in the efficient, dose-dependent knockdown of murine Gys1 mRNA. ASO#1 had an ED_{50} of 45.2 mg/kg (95% confidence interval (C.I.): 32.3-63.2 mg/kg) in quadriceps muscle, whereas ASO#2 was more potent, with an ED_{50} of 24.2 mg/kg (95% C.I.: 22.4-26.1 mg/kg) (Supplementary Figure 1A). Consistently, Gys1 ASO#2 achieved a robust reduction of GYS1 protein in skeletal muscle of wild type and Gaa$^{-/-}$ mice (Supplementary Figure 1B, 1C).

To determine if ASO-mediated Gys1 knockdown attenuates the pathological accumulation of glycogen in vivo, we dosed young Gaa$^{-/-}$ mice with the newly identified Gys1 ASO#1 and ASO#2. A starting age of one month was selected because at this time the Gaa$^{-/-}$ mice have not developed severe myopathy, yet, and the glycogen is only starting to accumulate in the skeletal muscle (Figure 1A). The one-month-old Gaa$^{-/-}$ mice (Cohort 1) were dosed via subcutaneous injection with 25 mg/kg Gys1 or control ASO, once a week for sixteen weeks (Figure 2A). Compared to control ASO, Gys1 ASO#1 and ASO#2 resulted in 83.64% (95% C.I.: 77.40-88.58) and 88.58% (95% C.I.: 83.61-92.06) Gys1 mRNA knockdown, respectively in quadriceps muscle (Figure 2B, Supplementary Table 1). Corroborating the reduction in mRNA expression, compared to control ASO, ASO#2 administration decreased GYS1 protein, measured by Western blot, in key Pompe tissues of Gaa$^{-/-}$ mice by 100%, 98%, and 100% in quadriceps, diaphragm, and heart, respectively. (Figure 2C, Supplementary Table 1).

To evaluate the impact of Gys1 knockdown on excessive glycogen accumulation in the muscle of the Pompe mice, we measured glycogen content by a colorimetric biochemical assay in the muscle

of the 20-week-old Gaa$^{-/-}$ mice following 16 weeks of Gys1 ASO treatment. ASO#1 resulted in a

38.06% reduction in glycogen content (0.29 μg/μg protein) in quadriceps muscle, and ASO#2

resulted in 57.05% reduction in glycogen (0.19 μg/μg protein) compared to control ASO (0.49

μg/μg protein) (Figure 2D, Supplementary Table 1). The superior efficacy of ASO#2 compared to

ASO#1 is consistent with its greater potency in knocking down Gys1 mRNA expression. Similarly,

the control ASO had no effect on Gys1 expression and glycogen content (Figure 2B-E). In

agreement with the biochemical measurements, semi-quantitative analysis of glycogen content in

histological sections using PAS staining confirmed that Gys1 ASOs ameliorated the aberrant

accumulation of glycogen in Gaa$^{-/-}$ quadriceps muscle. On a scale of 1 (no staining) to 4 (extensive

staining), an average score of 1.8 was assigned to the 4.5-month-old Gaa$^{-/-}$ mice treated with

control ASO, 1.4 to mice treated with ASO#1, and 1 to mice treated with ASO#2 (Figure 2E,

Supplementary Table 2).

In view of the enhanced efficacy in the early-treatment setting, we selected ASO#2 to further

investigate if Gys1 ASO treatment could revert the Pompe pathology in older Gaa$^{-/-}$ mice.

ASO-mediated Gys1 reduction partially reverses glycogen accumulation in skeletal muscle

of older Gaa$^{-/-}$ mice

To assess if therapeutic treatment with Gys1 ASO could reverse the glycogen accumulation and

myopathy in mature Pompe mice, we dosed older Gaa$^{-/-}$ mice with Gys1 ASO#2 starting at three

months of age (Cohort 2) (Figure 3A). Gys1 ASO#2 was administered by weekly subcutaneous

injections at 25 mg/kg for 16 weeks (Figure 3A). The rotarod motor performance assay showed

that the ASO#2-mediated knockdown of Gys1 expression resulted in a trend towards preservation

of motor function (Figure 3B), which otherwise progressively declines in Gaa$^{-/-}$ mice (Figure 1B,

3B). Indeed, the control ASO did not slow down the deterioration of muscle function in Pompe mice (Figure 3B). We sacrificed the animals at 6.5 months of age and studied the quadriceps muscle, diaphragm, and heart tissue from mice treated with Gys1 ASO#2. We measured a 64.71% (95% C.I.: 57.00-71.05) knockdown of mouse Gys1 mRNA and complete depletion of GYS1 protein in quadriceps muscle compared to control ASO (Figure 3C-D, Supplementary Table 1). Similarly, GYS1 protein was reduced by 80% in diaphragm and 70% in heart compared to mice treated with control ASO (Figure 3D, Supplementary Table 1). Of note, the expression of GYS1 was not significantly different in 6.5-month-old Pompe mice compared to wild type mice (Figure 3D). A biochemical assay demonstrated that the ASO#2-mediated knockdown of Gys1 expression was associated with a modest 7.79% (95% C.I.: 5.53-28.45) reduction in glycogen accumulation compared to control ASO in the quadriceps muscle of Gaa$^{-/-}$ mice (Figure 3E, Supplementary Table 1). In addition, semi-quantitative analysis of histological sections of quadriceps muscle using PAS staining confirmed a mild improvement of the aberrant glycogen accumulation following treatment with Gys1 ASO#2 (average score of 1.25 on a scale of 1 to 4) compared to control ASO (average score of 2.2) (Figure 3F, Supplementary Table 2).

ASO mediated Gys1 knockdown in combination with enzyme replacement therapy further reverses Pompe pathology, preserving motor function in older Gaa$^{-/-}$ mice

Substrate reduction therapy (SRT) using ASO to remove the source of new glycogen synthetized via GYS1 shows promise as a monotherapy, but the reversal of the myopathy in Gaa$^{-/-}$ mice was incomplete in 6.5-month-old animals. We reasoned that combining Gys1 ASO, which blocks the synthesis of new glycogen, with GAA enzyme replacement therapy (ERT) to simultaneously remove pre-existing glycogen in the skeletal muscle and heart of older Pompe mice, could provide

additive efficacy. Therefore, we dosed four-month-old Gaa$^{-/-}$ mice with Gys1 ASO#2 at 25 mg/kg

by weekly subcutaneous injections for six weeks, followed by a 10-week-long co-administration

phase where we combined ASO#2 and ERT, in the form of recombinant human alpha-acid

glucosidase (rhGAA) enzyme (Cohort 3) (Figure 4A). rhGAA was dosed intravenously every two

weeks (Figure 4A). Alternatively, Gaa$^{-/-}$ mice were dosed with either ASO#2 alone or ERT alone

(Figure 4A). Wild type mice and PBS-treated Gaa$^{-/-}$ mice were included in the study as controls.

The ASO#2+ERT combination treatment led to a stable increase in rotarod performance over the

4-month-long treatment, resulting in gains of 9, 8, 11 and 19% compared to baseline after 1, 2, 3,

and 4 months of treatment, respectively. On the other hand, ASO#2 alone increased the rotarod

performance in the first three months of treatment, but was followed by a slowdown (2, 10, 12 and

7% gain compared to baseline after 1, 2, 3, and 4 months of treatment, respectively). ERT alone

was unable to prevent the functional decline in the aging Pompe mice, resulting in a downward

change in rotarod performance over time (-2, -8, -5 and -8% compared to baseline after 1, 2, 3, and

4 months of treatment, respectively) (Figure 4B).

At the end of the study, the 8-month-old mice were sacrificed for further biochemical and IHC

analysis of the muscle tissue. Western blot analysis revealed that ASO#2 alone or ASO#2+ERT

resulted in the complete ablation of GYS1 protein expression in quadriceps muscle and heart

(Figure 4C). Similarly, minimal residual GYS1 protein was present in the diaphragm of the same

treatment groups (Figure 4C). On the other hand, Gaa$^{-/-}$ mice that received PBS or ERT alone had

robust GYS1 protein expression, comparable to wild type levels (Figure 4C). Importantly, the

systemic administration of Gys1 ASO#2, alone or in combination with ERT, was well tolerated in

Pompe mice with no significant changes in body or organ weights, and no significant elevation in

liver transaminases, blood urea nitrogen, or muscle creatine kinase in plasma compared to control

groups (Supplementary Figure 2).

The average glycogen content measured using the biochemical assay in quadriceps muscle of Gaa$^-$

$^{/-}$ mice was 24.85 (95% C.I.: 12.54-37.17), 10.98 (95% C.I.: 6.57-15.38), 4.84 (95% C.I. 1.61-

8.07), and 20.5 (95% C.I.: 12.46-28.53) µg glycogen per µg total protein in the PBS, ASO alone,

ASO+ERT, and ERT alone treatment groups, respectively (Figure 4D). Therefore, compared to

PBS controls, the biochemical glycogen assay revealed a 55.22% (95% C.I.: 20.29-74.84)

reduction in glycogen content in quadriceps muscle of Gaa$^{-/-}$ mice dosed with ASO#2 alone (Figure

4D, Supplementary Table 1). The ASO#2+ERT combination resulted in an even greater effect,

reducing glycogen levels by 83.48% (95% C.I.: 53.80-94.10) in quadriceps muscle (Figure 4D,

Supplementary Table 1). ERT alone did not significantly attenuate the glycogen accumulation,

leading to 15.78% (95% C.I.: 11.61-31.49) less glycogen in quadriceps muscle of Gaa$^{-/-}$ mice. ERT

was more efficacious in the heart, where it resulted in 65.82% (95% C.I.: 5.88-87.59) glycogen

reduction vs. PBS control when dosed alone, and in 74.93% (95% C.I.: 18.77-92.26) glycogen

reduction vs. PBS control when combined with Gys1 ASO#2 (Figure 4D, Supplementary Table

1). PAS staining of quadriceps muscle histological sections corroborated the biochemical assay

results, showing a marked reduction in glycogen content, especially in the ASO#2 group (semi-

quantitative score of 1.5 on a scale of 1 to 4, where 1 is no glycogen detected) and ASO#2+ERT

(score of 1), compared to PBS (score of 3.75) or ERT alone (score of 3) samples (Figure 4E,

Supplementary Table 2). Thus, the motor function improvement correlated well with the reduction

in muscle glycogen content.

Taken together, our data supports the hypothesis that blocking the synthesis of new glycogen while

concomitantly enhancing its clearance is beneficial in treating both the skeletal muscle and cardiac

abnormalities in Pompe disease (Figure 6).

ASO-mediated Gys1 knockdown with or without ERT, but not ERT alone, ameliorates the

autophagic buildup in the skeletal muscle of Gaa$^{-/-}$ mice

Defective autophagy is a major factor contributing to the muscle pathology in Pompe disease[7].

Consistent with the highly integrated functions of lysosomes and autophagosomes, it has been

shown that lysosomal dysfunction in Pompe disease leads to a block in autophagic flux and

accumulation of autophagic debris, particularly in muscle tissue[32]. It is important to note that the

Gaa$^{-/-}$ mouse model faithfully recapitulates the autophagic defects described in Pompe patients[30].

Incomplete autophagic flux can be assessed by measuring the aberrant accumulation of LC3 and

p62 proteins. Additionally, immunostaining of histological sections for lysosomal-associated

membrane protein 1 (LAMP1) provides a method to detect enlarged lysosomes[33,34]. We

investigated the efficacy of Gys1 ASO, alone or in combination with ERT (Cohort 3, Figure 4A),

in ameliorating the defective autophagy in Gaa$^{-/-}$ mice. Western blot analysis confirmed

accumulation of p62 and LC3-II proteins in quadriceps muscle and heart of 8-month-old Gaa$^{-/-}$

mice treated with PBS (Figure 5A). ERT alone alleviated the autophagic defect in the heart but not

significantly in the skeletal muscle, as indicated by the reduced levels of LC3-II, p62 and LAMP1

proteins in cardiac tissue with no major effect in quadriceps muscle (Figure 5A). On the other

hand, Gys1 ASO#2, alone or in combination with ERT, led to the almost complete normalization

of the p62, LC3-II and LAMP1 protein levels in quadriceps of Gaa$^{-/-}$ mice (Figure 5A). In the

heart, the Gys1 ASO contributed to a lesser extent to the reduction of the autophagic markers

(Figure 5A). In addition, immunostaining of histological sections confirmed that ASO#2+ERT

administration reduced LAMP1 protein levels in quadriceps muscle of $Gaa^{-/-}$ mice (Figure 5A,

5B), suggesting that the combination therapy could rescue the autophagy pathway dysfunction in

Pompe muscle after four months of treatment. The effect was more pronounced than in the Gys1

ASO#2-only group (Figure 5B).

In conclusion, our data indicate that the combination of Gys1 ASO and ERT might provide

therapeutic benefit by targeting various aspects of Pompe pathogenesis, namely glycogen

accumulation and autophagy defects.

Systemic Gys1 knockdown does not mitigate Pompe pathology in the central nervous system

Pompe disease is a complex multi-systemic condition with growing evidence of central nervous

system (CNS) involvement[35]. Indeed, we found remarkable abnormalities in the spinal cord of

$Gaa^{-/-}$ mice at 8 months of age (Supplementary Figure 3). The pathologic changes were visible in

histological sections of the spinal cord, where Nissl staining (toluidine blue) revealed the presence

of swollen cell bodies, activated glial cells, and centronucleated cells (Supplementary Figure 3).

Systemically delivered Gys1 ASO#2 did not ameliorate the Pompe CNS phenotype of $Gaa^{-/-}$ mice

(Supplementary Figure 3), consistent with the inability of ASOs to cross the blood-brain barrier.

DISCUSSION

Enzyme replacement therapy, currently the standard of care for patients with PD, has been shown

to be beneficial, but its efficacy is hindered by some limitations, especially in treating the skeletal

muscle manifestations of PD. Early administration of ERT improves cardiac hypertrophy and

motor development, extending the chances of survival in patients with infantile-onset disease

(IOPD)[6,36]. Patients with the late-onset form (LOPD) have early increase and stabilization of their

pulmonary and motor function, however after reaching a plateau there is continued decline in their

motor function such as the 6 Minute Walk Test[8,9]. Progressive muscle weakness thus continues to

be a problem for long-term survivors of PD[5,6]. An explanation is provided by studies that have

shown that the delivery of the therapeutic enzyme to skeletal muscle is insufficient to reduce the

glycogen content and eliminate the autophagic buildup[6]. Recently, a study using a new adjunct

therapy based on the enzyme and chaperone ATB200/AT222 combination showed enhanced

efficacy over the currently approved ERT in Gaa$^{-/-}$ mice. ATB200/AT222 was able to normalize

muscle glycogen content and reduced the levels of lysosomal and autophagosomal markers in the

muscle of Gaa$^{-/-}$ mice to wild type levels[7,18]. Clinical trials of AT2221/ATB200 have been

completed and this approved treatment now offers an alternative effective therapeutic option for

Pompe disease patients.

Douillard-Guilloux et al. (2008) demonstrated that intramuscular injection of AAV-1 expressing

shRNA targeted to Gys1 reduces glycogen accumulation in the gastrocnemius of newborn Gaa$^{-/-}$

mice[19]. The same research group also analyzed the effect of a complete genetic ablation of

glycogen synthesis in a new Gaa/Gys1 double knock-out (KO) mouse model[17]. The Gaa/Gys1 KO

mouse exhibited a profound reduction in cardiac and skeletal muscle glycogen, a significant

decrease in lysosomal swelling and autophagic buildup, and complete correction of

cardiomegaly[17]. Muscle atrophy, observed in 11-month-old single Gaa KO mice, was less

pronounced in the Gaa/Gys1 double KO mice, resulting in improved exercise capacity[17]. This data

demonstrated that long-term suppression of muscle glycogen synthesis is well tolerated and leads

to a significant improvement in Pompe pathology. However, no approved small molecule

inhibitors of muscle GYS1 are currently clinically available, underscoring the challenges of

finding specific compounds that can preserve the activity of the hepatic GYS2 enzyme.

ASO technology has emerged as a highly specific therapeutic option for a variety of neuromuscular

diseases. ASOs that modulate exon splicing of the dystrophin (DMD) gene can reduce muscle

degeneration in Duchenne's muscular dystrophy[37-39,11,40,41]. An ASO that disrupts spliceosome

action on exon 7 of the survival motor neuron 1 (SMN1) gene and restores the gene to functioning

length has been approved for treatment of spinal muscular atrophy[42,43]. Recently, an ASO with

RNase H1 mechanism has been approved by the FDA for amyotrophic lateral sclerosis (ALS)

patients with SOD1 mutations[41,42]. Several studies have attempted to use ASOs to mitigate the

disease outcomes by improving or restoring normal GAA synthesis[6]. Van der Wal et al. (2017)

studied the common IVS132-13T>G GAA mutation and located sequences where an ASO could

block splicing machinery and prevent exclusion of exon 2 in the final transcript, leading to

increased GAA expression[44,45]. Goina et al. (2017) performed a similar exon 2 splicing modulation

with an ASO and successfully reduced glycogen accumulation in patient-derived myotubes[46].

Other ASO studies have sought to reduce glycogen by interfering with GYS1 expression. Clayton

et al. (2014) demonstrated that treating Pompe mice with a phosphorodiamidate morpholino

oligonucleotide (PMO) designed to induce exon 6 skipping in the murine Gys1 mRNA resulted in

the degradation of messenger RNA due to frame shift[16]. However, to facilitate PMO delivery to

muscle via systemic administration, the PMO was conjugated with a cell penetrating peptide (GS-

PPMO), and the development of PPMO drugs for human use is significantly challenged by their

potential toxicity, as observed in primate studies[46]. This toxicity is likely due to the cationic nature

of the peptide at the doses required for efficacious exon skipping in the primate musculature[16,47].

Benefiting from advancements in ASO technology, here we identified a potent ASO that safely

knocks down mouse Gys1 expression by RNase H1 mechanism in mice. Using both prevention

and reversal treatment modalities, we demonstrated the potential of the Gys1 ASO approach to

serve as efficacious SRT in the Gaa$^{-/-}$ mouse model of Pompe disease.

The Gaa$^{-/-}$ Pompe mice start to accumulate glycogen in skeletal muscle as early as three weeks of

age but develop a progressive defect in motor functions from six months of age. We have attempted

to address the importance of timing in two separate cohorts, starting the ASO treatment at one-

and three-month-old mice respectively. Taken together, our data indicate that long-term

elimination of muscle glycogen synthesis leads to a significant reduction of glycogen accumulation

and autophagic buildup, as well as a tendency towards functional improvement in severely affected

older Pompe mice. Interestingly, we observed superior activity of Gys1 ASO in the skeletal muscle

of Pompe mice compared to wild type, healthy mice (Supplementary Figure 1). We hypothesize

this phenomenon being the result of increased permeability to ASO of the damaged muscle

combined with facilitated escape of the ASO from defective endosomal compartments upon

cellular uptake.

Despite the positive results obtained by knocking down Gys1 expression with ASO, there was still

a residual amount of glycogen present in the skeletal muscle of older Gaa$^{-/-}$ mice, and the muscle

structure abnormality was not significantly improved upon treatment with ASO alone

(Supplementary Table 2). This could be due to compensatory mechanisms, or very low levels of

residual GYS1 enzymatic activity. To further improve the efficacy of our Gys1 ASO-based SRT

approach both in the skeletal muscle and heart, we decided to combine the ASO treatment with

ERT. Our data shows that the combination of Gys1 ASO with rhGAA results in enhanced

reduction of glycogen content and reversal of the autophagy defect in the skeletal muscle of old

Gaa$^{-/-}$ mice. The molecular changes induced by the ASO and ERT treatment led to the preservation

of motor function in the aging Pompe mice. Recently Ullman et al. (2024)[48] recently reported their

preclinical studies of MZ-101, a selective and noncompetitive inhibitor of GYS1, reduced

glycogen accumulation, similar to that of ERT alone in Pompe mouse models.   MZ-101 when

utilized in combination with ERT however had an additive effect of normalizing glycogen

concentrations in the muscles. These data from prior studies and our promising data suggests

that substrate reduction therapy with GYS1 inhibition may be a promising therapeutic approach

for Pompe disease. Recently, Maze Therapeutics announced positive phase 1 study results from a

clinical trial of the small molecule GYS1 inhibitor MZE001 (ClinicalTrials.gov Identifier:

NCT05249621). In the study the healthy volunteers demonstrated exposure-dependent

reductions in blood mononuclear cell glycogen across dose levels 10 days after administration. It

is also expected that GYS1 ASOs has the potential to be used as monotherapy or in combination

with ERT in future clinical trials in Pompe patients. The results from the clinical trial further

support the validity and safety of GYS1 inhibition as a potential therapeutic approach for Pompe

disease.

MATERIALS AND METHODS

Identification of antisense oligonucleotides (ASOs) targeting mouse Gys1

All ASOs used in the studies described here were 16 nucleotides in length with a phosphorothioate

(PS) backbone and three 2'-constrained ethyl (cEt)-modified nucleotides at both ends (3-10-3

gapmer configuration). A well-characterized control ASO which does not hybridize to any mouse

mRNA sequence was included in the experiments. The oligonucleotides were synthesized and

purified as previously described[49]. ASOs targeting murine Gys1 were designed and initially

screened for activity in B16-F10 cells via electroporation. The most active ASOs were

subsequently evaluated in wild type mice to identify ASOs that were safe and active in vivo. A

dose-response confirmation experiment for the two lead ASOs, ASO#1 and ASO#2, was

performed in wild type mice by subcutaneous injection twice a week for 5.5 weeks at 12.5, 25, 50,

and 100 mg/kg total weekly doses (shown in Supplementary Figure 1A). All ASOs were

formulated in phosphate buffered saline (PBS) and were administered to mice by subcutaneous

injection. The mouse ASO screening studies were conducted under protocols approved by the

Institutional Animal Care and Use Committees at Ionis Pharmaceuticals.

Characterization of the Gaa$^{-/-}$ animal model of Pompe disease

The Gaa$^{-/-}$ Pompe mouse model was a gift of Dr. Nina Raben[50]. The mouse model is a constitutive

homozygote knockout of the murine acid alpha-glucosidase (Gaa) gene. One-month-old Gaa$^{-/-}$

mice, which show only the initial signs of tissue glycogen accumulation without severe myopathy,

were used in prevention studies, whereas three-month-old Gaa$^{-/-}$ mice, which show marked

glycogen accumulation and motor function impairment, were used in reversal studies. All the

intervention studies performed in either the Gaa$^{-/-}$ mouse model or wild type C57BL/6 control mice

were conducted in accordance with the Institutional Animal Care and Use Committee at the

University of California, Irvine (UCI), under protocol IACUC AUP-19-075, and were consistent

with Federal guidelines. The mice were housed at UCI animal facility that maintained a constant

temperature (22°C) and controlled light and dark cycles (12:12). The overall condition of the mice

was monitored daily throughout the experimental process to assess general health and mitigate any

pain and suffering.

Administration of ASOs to Gaa$^{-/-}$ mice

Gaa$^{-/-}$ or wild type control mice were dosed with ASOs once a week at 25 mg/kg via subcutaneous

injection. For the early-treatment study (Cohort 1) described in Figure 2, one-month-old Gaa$^{-/-}$

mice were randomly divided into groups for treatment with different Gys1 ASOs, control ASO, or

vehicle (PBS), n=12-17 mice per group. For the reversal study (Cohort 2) described in Figure 3,

three-month-old Gaa$^{-/-}$ mice were randomly divided into three groups for treatment with Gys1

ASO#2 (n=15), control ASO (n=10), or vehicle (PBS) (n=10). The mice received a total of 16

weekly ASO doses (16 weeks duration) and were sacrificed 48 hours after the last ASO dose. Body

weight changes, and the weight of liver, kidney, and spleen were recorded to exclude potential

toxicity of the ASOs in the context of the long dosing paradigm in diseased Gaa$^{-/-}$ mice.

Adjunct treatment using Gys1 ASO in combination with ERT in Gaa$^{-/-}$ mice

Four-month-old Gaa$^{-/-}$ mice were dosed with Gys1 ASO#2 or control ASO by subcutaneous

injection once a week at 25 mg/kg for six weeks. Starting from week 7 the mice additionally

received 20 mg/kg recombinant human GAA (rhGAA) (Genzyme Corporation, Cambridge, MA;

Cat.: NDC 58468-0160-1) via intravenous injection once every two weeks. The antihistaminic

agent diphenhydramine hydrochloride was administered to the mice via intraperitoneal injection

15 minutes before receiving the human enzyme, starting from the second rhGAA administration.

The weekly subcutaneous ASO administration continued throughout the ERT phase for a total of

16 doses. The mice were euthanized at 8 months of age, after having been treated with Gys1 ASO

for 4 months, the last two of which overlapped with the ERT; n=5 per treatment group.

Accelerating rotarod performance test

Muscle function was evaluated using a rotarod apparatus (Rotamex-5, Columbus Instruments,

Columbus, OH). Briefly, mice were placed on a rotarod whose speed was set to progressively

increase from 4 to 40 rotations per minute (rpm) in 300 seconds. The latency to fall from the

rotating rod was recorded as a measure of muscle strength. Assessments were done on three

consecutive days, three to five trials each day with at least two-minute rest intervals between the

trials. The mean latency time to fall off the rotarod from three independent trials from the last two

days was used for analysis.

Biochemical analysis of serum from mice for toxicity studies

Blood samples were collected from mice via cardiac puncture under deep anesthesia using

isoflurane inhalation. Immediately after collection, approximately 300-500 μL of whole blood

were placed into serum separator tubes (Greiner Bio-One Minicollect tubes; Cat.: 450472).

Samples were allowed to clot for at least 20 minutes at room temperature, and then centrifuged at

3,000 rpm for 15 minutes. Serum was transferred to fresh tubes and stored at -80°C before being

used for clinical biochemistry evaluation at IDEXX Laboratories (IDEXX Laboratories Inc.,

Maine, USA). Parameters measured include alanine aminotransferase, aspartate aminotransferase,

alkaline phosphatase, blood urea nitrogen, creatine kinase.

Gene expression analysis using quantitative real-time PCR

Tissue samples were isolated from mice and snap frozen in liquid nitrogen. Frozen tissues were

homogenized in TRIzol reagent (Thermo Scientific, Waltham, MA; Cat.: 15596026), and total

RNA was extracted according to the manufacturer's instructions. Real-time qPCR analysis was

performed using the following reagents: mouse Gys1 (Taqman™ assay ID: Mm01962575_s1),

mouse Gapdh (Taqman™ assay ID: Mm 99999915_g1, housekeeping gene). In the initial

assessment of ASO activity in wild type mice (Supplementary Figure 1A) real-time qPCR analysis

was performed using the following primer-probe sets: mouse Gys1 (forward primer: 5'-

TGATGAAGAGAGCCATCTTTGC-3'; reverse primer: 5'-AGGAGTCGTCCAGCATGTTGT-

3'; probe: 5'-ACTCAGCGGCAGTCTTTCCCACCA-3'), mouse cyclophilin A (forward primer:

5'-TCGCCGCTTGCTGCA-3'; reverse primer: 5'-ATCGGCCGTGATGTCGA-3'; probe: 5'-

CCATGGTCAACCCCACCGTGTTC-3'). Reactions were conducted in triplicate on a

QuantStudio7 real-time PCR system (Thermo Fisher Scientific, Waltham, MA), following the

manufacturer's instructions.

Biochemical measurement of tissue glycogen content

Mouse tissues were collected, snap frozen in liquid nitrogen upon collection, and stored at -80ºC.

Approximately 10 mg of frozen quadriceps muscle were rinsed in 300 μL of ice-cold PBS, then

homogenized in 200 μL of double-distilled water using a Wheaton dounce tissue grinder (DWK

Life Sciences, Rockwood, TN) as follows: 10-20 passes on loose pestle followed by 10-20 passes

on tight pestle, keeping the tissue grinder on ice. The tissue lysates were then heated at 100°C for

10 minutes, then centrifuged at 15,000 g at 4°C for 10 minutes. Total protein concentration in the

supernatant was determined using the Micro BCA protein assay kit (Thermo Scientific, Waltham,

MA; Cat.: 23235). In the same supernatant, glycogen concentration was measured using a

fluorometric enzymatic assay (Sigma-Aldrich, St. Louis, MO; Cat.: MAK016), according to the

manufacturer's instructions. The assay produces a fluorometric (λex = 535/λem = 587 nm) product

proportional to the glycogen present in the sample. The fluorescent signal was analyzed using the

BioTek Gen5 software (Agilent Technologies, La Jolla, CA), and normalized by the total protein

content of the sample.

Western blot analysis of autophagy markers

Approximately 30 mg of frozen tissue was homogenized in 500 µL of RIPA buffer (Sigma-

Aldrich, St. Louis, MO; Cat.: R0278) supplemented with a proteinase inhibitor cocktail (Sigma-

Aldrich, St. Louis, MO; Cat.: P8340) using a handheld electric homogenizer (Qsonica, Newtown,

CT; Cat.: XL-2000). The tissue homogenate was then agitated on a Benchmark Roto-Mini Plus

(Benchmark Scientific, Sayreville, NJ) for 2 hours at 4°C. Lastly, the tissue lysate was centrifuged

at 20,000 g at 4°C for 30 minutes, and the supernatant was collected for analysis. The sample

protein concentration was measured using the Micro BCA protein assay kit (Thermo Fisher

Scientific, Waltham, MA; Cat.: 23235). 20 µg of total protein lysate was resolved by

electrophoresis on Bis-Tris 4-12% NuPAGE polyacrylamide gels using the Novex Mini Cell

system (Invitrogen/Thermo Fisher Scientific, Waltham, MA). The proteins were then transferred

to a PVDF membrane, which was later blocked with 1% bovine serum albumin (BSA) in Tris-

buffered saline containing 0.1% Tween 20 (TBS-T). The primary antibodies used in our studies

were: anti-GYS1 (Abcam, Waltham, MA; Cat.: ab 40810; 1:5,000 dilution), anti-LC3B-I/II

(Abcam, Waltham, MA; Cat.: ab192890; 1:5,000 dilution), anti-p62 (Abcam, Waltham, MA; Cat.:

ab56416; 1:20,000 dilution), anti-LAMP1 (Abcam, Waltham, MA; Cat.: ab27170; 1:2,000

dilution), anti-GAPDH (Abcam, Waltham, MA; Cat.: ab18160; 1:10,000 dilution; served as

loading control), anti-β-ACTIN (Abcam, Waltham, MA; Cat.: ab8227; 1:5,000 dilution; served as

alternative loading control). Horseradish peroxidase (HRP)-linked goat anti-mouse or goat anti-

rabbit secondary antibodies were used (1:5,000 dilution). Densitometric quantification was

performed using ImageJ software (National Institutes of Health, Bethesda, MD).

Periodic acid-Schiff Histological analysis

Periodic acid-Schiff (PAS) staining was performed to detect glycogen in histological tissue

sections. Briefly, the quadriceps muscle was dissected and fixed in a 4% neutral buffered formalin

solution at room temperature for 20 hours. The fixed tissue was then rinsed in PBS and transferred

to a 70% ethanol solution before being embedded in paraffin blocks and sectioned. The PAS

staining was performed on 8μm-thick sections by the Pathology Department at UCI Medical

Center according to protocols described previously[49]. Images of the histological slices were

captured using a Keyence BZ-810 microscope (Keyence, Itasca, IL). In PAS-stained tissue

sections, glycogen appears fuchsia-colored under light microscopy. For quantification of the PAS

staining, the histological sections of quadriceps muscle from 4-5 animals per treatment group were

analyzed, and a semi-quantitative score from 1 (no staining) to 4 (extensive staining) was assigned

to each section. The scoring was performed independently by two different technicians to ensure

accurate assessment. To quantify the number of myofibers with centralized nuclei, a minimum of

300 myofibers per animal were manually evaluated in the PAS-stained sections of quadriceps

muscle (n=4-5 mice per each group). The data is presented as the percentage of myofibers with a

centralized nucleus.

Immunohistochemistry (IHC) studies

For lysosomal analysis, the expression of lysosomal associated membrane protein 1 (LAMP1)

was assessed using IHC in tissue sections of quadriceps muscle embedded in frozen section

medium (Fisher Scientific, Hampton, NH; Cat.: 22-046-511), as described previously[51]. Briefly,

8µm-thick cryosections (10 sections per each animal, 5 animals per treatment group) were fixed

in 4% paraformaldehyde (PFA), blocked in donkey serum, and stained with an anti-LAMP1

primary antibody (Abcam, Waltham, MA; Cat.: ab24170; 1:500 dilution) at 4°C overnight. The

sections were then washed three times with PBS and incubated with fluorescein-conjugated

secondary antibody (Thermo Fisher Scientific, Waltham, MA; Cat.: A10042; 1:500 dilution) for

1 hour at room temperature, then washed and mounted with DAPI-containing mounting media

(Vectashield, Vector Laboratories, Newark, CA; Cat.: H-1200-10). The slides were imaged on a

Zeiss LSM 900 confocal microscope (Carl Zeiss Microscopy, White Plains, NY). For the

quantification of the LAMP1 staining, the percentage of LAMP1-positive myofibers was

calculated and compared across treatment groups.

Nissl (toluidine blue) staining was used to study the morphology and pathology of neural tissue.

The spinal cord was harvested from mice, fixed in 4% paraformaldehyde (PFA) for 24 hours, and

then incubated in a solution containing 30% sucrose in PBS overnight, embedded in optimal

cutting temperature compound (OCT compound), and cryo-sectioned. Subsequently, the 10-

micron cryosections were stained with 1% toluidine blue containing 1% borate solution as

previously described[52,53]. Briefly, the slices were placed in Toluidine blue solution for 30 seconds

and rinsed in triple distilled water. Stained slices were examined using light microscopy.

Statistical analysis

Statistical analysis was performed using GraphPad Prism software (GraphPad Software, Boston,

MA), or RStudio software (RStudio, Boston, MA) in the case of Figure 1B. Unpaired t-test, one-

way ANOVA, or two-way ANOVA were used, as specified in the respective figure legends. p-

values lower than 0.05 were considered statistically significant; *: $p<0.05$, **: $p<0.01$, ***: $p <$

0.005, and ****: $p<0.0001$, ns= not significant. Error bars indicate the standard error of the mean

(SEM).

Legends

Figure 1. Gaa$^{-/-}$ mice show progressive accumulation of glycogen in skeletal muscle followed

by deterioration of muscle function.

A. Glycogen content in quadriceps from Gaa$^{-/-}$ mice at 1, 2, and 3 months (mo) of age compared

to 1-month-old wild type (WT) C57BL/6 mice. n=5 per group, as indicated in parenthesis on the

x-axis. Statistical analysis was performed using unpaired t-test in GraphPad Prism software. **B.**

Graph illustrates the time that the mice spent on an accelerating rotarod before falling, to assess

muscle strength as the animals age from 2 to 14 months. The mean decline in motor function is

represented by the blue line within the shaded grey area, which indicates the SEM range. The

number of mice in each measurement group is indicated at the bottom of the graph (n). Statistical

analysis was performed using one-way ANOVA in GraphPad Prism software. *: $p<0.05$, **: $p <$

0.01, ***: $p < 0.005$, ****: $p < 0.001$.

Figure 2. Early treatment of Gaa$^{-/-}$ mice with Gys1 ASOs lowers target mRNA and protein

expression, leading to reduced glycogen accumulation in skeletal muscle.

A. Schematic illustration of the early treatment paradigm with Gys1 ASOs in Gaa$^{-/-}$ mice from 1

to 4.5 months of age. WT: wild type. **B.** qPCR analysis of Gys1 mRNA in quadriceps muscle. **C.**

Representative images of Western blot analysis of GYS1 protein in quadriceps, diaphragm, and

heart. The number of samples in each group is listed in parentheses on the x-axis. **D.** Glycogen

content measured using a biochemical assay in quadriceps muscle from Gaa$^{-/-}$ mice dosed with

either PBS, control ASO (Ctrl ASO), or two different Gys1 ASOs. The number of samples in each

group is listed in parentheses on the x-axis. **E.** Periodic acid-Schiff (PAS) staining of histological

sections of quadriceps muscle. Scale bars: 50 μm.

Statistical analysis was performed using unpaired t-test in GraphPad Prism software. ***: p <

0.005, ****: p < 0.0001.

Figure 3. Therapeutic treatment with Gys1 ASO in Gaa$^{-/-}$ mice partially reverses glycogen

accumulation in skeletal muscle and preserves muscle strength.

A. Schematic illustration of the reversal treatment paradigm with Gys1 ASO#2 in Gaa$^{-/-}$ mice from

3 to 6.5 months of age. WT: wild type. **B.** Graph illustrates the time that mice spent on an

accelerating rotarod before falling, to assess muscle strength. **C.** qPCR analysis of Gys1 mRNA.

The number of samples in each group is listed in parentheses on the x-axis. **D.** Representative

images of Western blot analysis of GYS1 protein in quadriceps, diaphragm, and heart. GAPDH

was used as loading control. **E.** Glycogen content measured using a biochemical assay in

quadriceps muscle from Gaa$^{-/-}$ mice dosed with either PBS, a control ASO (Ctrl ASO), or Gys1

ASO. The number of samples in each group is listed in parentheses on the x-axis. **F.** Representative

images from PAS staining of histological sections of quadriceps muscle. Scale bars: 50 μm.

Statistical analysis was performed using unpaired t-test in GraphPad Prism software. *: p<0.05,

***: p < 0.005, ****: p < 0.001, ns: not significant.

Figure 4. Combination treatment with Gys1 ASO and ERT reduces glycogen accumulation

and improves motor function in Gaa$^{-/-}$ mice.

A. Schematic illustration of the study where 4-month-old Gaa$^{-/-}$ mice were dosed with Gys1

ASO#2 and ERT in combination. WT: wild type. **B.** Graph shows the results of a rotarod test to

assess muscle strength. The time that the mice spent on a rotating rod was recorded monthly over

the three months of treatment and expressed as percent change versus baseline measurements

(collected before the start of dosing) for each group; n=5 per group. Statistical analysis in B was

performed by two-way ANOVA using multiple comparisons in GraphPad Prism software. ****:

$p < 0.001$ between the $Gaa^{-/-}$ mice that received PBS and Gys1 ASO#2+ERT. **C.** Representative

images of Western blot analysis of GYS1 protein in quadriceps muscle, diaphragm, and heart.

GAPDH was used as loading control. **D.** Glycogen content measured using a biochemical assay in

quadriceps muscle and heart from wild type or $Gaa^{-/-}$ mice dosed with either PBS, Gys1 ASO#2,

Gys1 ASO#2+ERT, or ERT alone (n=5 per group).

Statistical analysis in C and D was performed using unpaired t-test. *: $p<0.05$, **: $p < 0.01$, ***:

$p < 0.005$, ns: not significant.

Figure 5. Normalization of autophagic flux in skeletal muscle of Pompe mice after treatment

with Gys1 ASO, alone or in combination with ERT.

A. Western blot analysis of LC3-II/LC3-I, p62, and LAMP1 proteins in quadriceps muscle and

heart from mice in cohort 3. GAPDH was used as loading control. The graphs illustrate the

quantification of Western blots from three independent experiments. WT: wild type. **B.**

Representative images of LAMP1 immunostaining in histological sections of quadriceps muscle

from $Gaa^{-/-}$ mice treated with Gys1 ASO#2, alone or in combination with ERT. The graph shows

the quantification of the stain as percentage of LAMP1-positive cells. LAMP1: lysosomal-

associated membrane protein 1. Statistical analysis was performed using unpaired t-test. *: $p<0.05$,

**: $p < 0.01$, ***: $p < 0.005$, ns: not significant.

Figure 6: **Model of pharmacological intervention in Pompe disease combining GYS1 ASO-**

based SRT and ERT.

We propose an adjunct therapeutic approach for Pompe disease based on the reduction of the

synthesis of new glycogen via GYS1 ASO-mediated SRT, combined with the ERT-mediated

clearance of pre-existing glycogen that has accumulated in the lysosome because of GAA

inactivation. Such combinatorial strategy would result in greater clinical benefit for Pompe

patients, especially in skeletal muscle, where ERT alone is not remarkably effective. SRT:

substrate reduction therapy; ERT: enzyme replacement therapy; GYS1: glycogen synthase 1;

ASO: antisense oligonucleotide; GAA: glucosidase, alpha acid; rhGAA: recombinant human

GAA.

Supplementary Figure 1: Gys1 ASO#1 and ASO#2 knock down mouse Gys1 mRNA and protein in wild type mice.

A. Knockdown of mouse Gys1 mRNA measured by qPCR in quadriceps muscle of wild type mice treated with ASO#1 and ASO#2 by subcutaneous injection at the indicated total weekly doses; n=4 per group. **B.** Western blot analysis of GYS1 protein in gastrocnemius muscle of wild type mice treated via subcutaneous injection with ASO#2 for 6 weeks at the indicated total weekly doses; n=4 per group. The graph shows the quantification of GYS1 protein expression relative to GAPDH (loading control) from three independent experiments. The GYS1 expression level after treatment with control ASO (Ctrl ASO) at 100 mg/kg total weekly dose was used as reference and was therefore set to 1. **C.** Western blot analysis of GYS1 protein in quadriceps muscle of Gaa$^{-/-}$ mice treated via subcutaneous injection with ASO#2 for 6 weeks at the indicated weekly doses; n=3 per group. The PBS control group was used as reference to determine normal GYS1 levels and was therefore set to 1. Representative blots from 3 sets of experiments are shown.

Supplementary Figure 2. Schematic design of the treatment strategy and toxicity studies using Gys1 ASO alone or in combination with ERT in Gaa-/- mice. Gys1 ASO and ERT are well tolerated in Gaa$^{-/-}$ mice.

A. Schematic illustration of the study used to assess the safety of the Gys1 ASOs in vivo. **B.** Change in body weight of wild type and Gaa$^{-/-}$ mice dosed with either PBS, Gys1 ASO#2 alone, or Gys1 ASO#2 in combination with ERT. The mice were dosed from 4 to 8 months of age, and their body weights were recorded once every four weeks during the four-month study. **C.** Terminal organ weight to body weight ratio normalized to wild type control group (set to 1). All the treatments were well tolerated since the ratios are within two-fold of wild type. **D.** Serum toxicology analysis performed at the end of the study shows no statistically significant change in the treatment groups compared to control Gaa$^{-/-}$ mice dosed with PBS. The number of samples in each group is listed in parentheses on the x-axis. ALT: alanine transaminase (values above 100 U/L are considered a sign of liver toxicity); AST: aspartate transaminase (values above 250-300 U/L are considered a sign of liver toxicity); BUN: blood urea nitrogen (values above 30 mmol/L are considered a sign of kidney toxicity); CK: creatine kinase (values above 1,500 U/L are considered a sign of muscle toxicity).

Supplementary Figure 3. Histological analysis of spinal cord in 8-month-old Gaa$^{-/-}$ mice with systemically delivered Gys1 ASO. No effect of systemically delivered Gys1 ASO in CNS. Nissl (toluidine blue) staining of histological sections of spinal cord from 8-months-old Gaa$^{-/-}$ mice dosed by subcutaneous injection with 25 mg/kg of either control ASO or Gys1 ASO#2 once a week for 16 weeks (16 total doses, 3.5 months of treatment, Cohort 2). Solid arrows indicate swollen cell bodies in the cervical ventral horn; dashed arrows indicate activated glial cells; asterisks indicate eccentric nuclei.

Supplementary Table 1. Summary of percent reduction over control ASO or PBS of Gys1 mRNA, glycogen content, and GYS1 protein in Gaa$^{-/-}$ mice treated with ASO, ASO+ERT, or ERT alone.

Mice from three separate cohorts were dosed with the indicated treatment, starting at 1, 3, or 4 months of age, and ending at 4.5, 6.5, or 8 months of age, respectively. The data is reported as percent reduction over control ASO or PBS of Gys1 mRNA, glycogen content, and GYS1 protein in Gaa$^{-/-}$ mice treated with ASO, ASO+ERT, or ERT alone

		% reduction					
		Gys1 mRNA (±95% C.I.)	GYS1 protein			Glycogen content (±95% C.I.)	
Mouse cohort	**Treatment**	**Quadriceps**	**Quadriceps**	**Diaphragm**	**Heart**	**Quadriceps**	**Heart**
1-month-old Gaa$^{-/-}$ (vs. Control ASO)	ASO#1 (n=12)	83.64 (77.40 - 88.16)	98	85	90	38.06 (14.50-61.07)	N.M.
	ASO#2 (n=12)	88.58 (83.61 - 92.06)	100	98	100	57.05 (35.57-71.36)	N.M.
3-month-old Gaa$^{-/-}$ (vs. Control ASO)	ASO#2 (n=10)	64.71 (57.00 - 71.05)	100	80	70	7.97 (5.53-28.45)	N.M.
4-month-old Gaa$^{-/-}$ (vs. PBS)	ASO#2 (n=5)	N.M.	100	90	100	55.22 (20.29-74.84)	14.03 (28.58-47.22)
	ASO#2+ERT (n=5)	N.M.	100	85	90	83.48 (53.80-94.10)	65.82 (5.88-87.59)
	ERT (n=5)	N.M.	0	0	0	15.78 (11.61-31.49)	74.93 (18.77-92.26)

697   N.M.: not measured; C.I: confidence interval.

698 **Supplementary Table 2. Summary of histological analyses.**

699 Semi-quantitative analysis of PAS staining, and percentage count of myofibers with a centralized

700 nucleus.

Mouse cohort	Genotype	Treatment	Quadriceps PAS score (1 to 4)	% Myofibers with centralized nucleus
1-month-old	$Gaa^{+/+}$	WT (n=5)	1	0
	$Gaa^{-/-}$	PBS (n=5)	2.2	8.4
		Ctrl ASO (n=5)	1.8	5.4
		ASO#1 (n=5)	1.4	14.1
		ASO#2 (n=5)	1	6.1
3-month-old	$Gaa^{+/+}$	WT (n=5)	1	0.7
	$Gaa^{-/-}$	PBS (n=5)	3.2	25.7
		Ctrl ASO (n=5)	2.2	19.6
		ASO#2 (n=5)	1.25	23.4
4-month-old	$Gaa^{+/+}$	WT (n=4)	1	1.4
	$Gaa^{-/-}$	PBS (n=4)	3.75	26.9
		ASO#2 (n=4)	1.5	25.2
		ASO#2+ERT (n=4)	1	16
		ERT (n=4)	3	17.9

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A

Quadriceps

Glycogen (μg/μg protein)

0.8

0.6 — ***

0.4 — *

0.2 — ***

0.0

1mo (5)   1mo (5)   2mo (5)   3mo (5)

WT   Gaa⁻/⁻

B

Rotarod

*   **   ****

Time before falling (seconds)

Age in months:	2	3	4	5	6	7	8	9	10	11	12	13	14
n:	11	20	20	20	30	30	27	10	17	9	10	9	11

A

Cohort 1: 1-month-old Gaa$^{-/-}$ study design

ASO#1,2

25 mg/kg every week
16 weeks
subcutaneous injection

WT, Gaa$^{-/-}$

Birth | 1 | 4.5 (months old)

B Quadriceps

mGys1 mRNA
(% WT control)

150%
100%
50%
0%

**** ****

WT | Ctrl ASO | ASO#1 (12) | ASO#2 (12)

Gaa$^{-/-}$

C Quadriceps | Diaphragm | Heart

WT | PBS | Ctrl ASO | ASO#1 | ASO#2

GYS1
β-Actin

Gaa$^{-/-}$

D Quadriceps

Glycogen
(µg/µg protein)

0.8
0.6
0.4
0.2
0.0

*** ****

PBS (12) | Ctrl ASO (17) | ASO#1 (12) | ASO#2 (12)

Gaa$^{-/-}$

E WT | Gaa$^{-/-}$

PBS | Ctrl ASO

ASO#1 | ASO#2

Quadriceps

Cohort 2: 3-month-old Gaa⁻/⁻ study design

ASO#2

25 mg/kg every week
16 weeks
subcutaneous injection

WT, Gaa⁻/⁻

Birth   3   6.5 (months old)

Time before falling (seconds)

ns   ns   *   ns

Month of Age   3   6.5   3   6.5   3   6.5   3   6.5
                 WT (10)   PBS (10)   Ctrl ASO (10)   ASO#2 (15)

Quadriceps

mGys1 mRNA (% PBS control)

****

PBS (10)   Ctrl ASO (10)   ASO#2 (10)

D

| Quadriceps | Diaphragm | Heart |

GYS1
GAPDH

WT PBS Ctrl ASO ASO#2   WT PBS Ctrl ASO ASO#2   WT PBS Ctrl ASO ASO#2

Gaa⁻/⁻   Gaa⁻/⁻   Gaa⁻/⁻

E

Quadriceps

Glycogen (µg/µg protein)

*
ns

PBS (10)   Ctrl ASO (10)   ASO#2 (10)

Gaa⁻/⁻

WT   Gaa⁻/⁻

PBS   Ctrl ASO   ASO#2

Quadriceps

A Cohort 3: 5-month-old Gaa$^{-/-}$ adjunct treatment study design

ERT
20 mg/kg every 2 weeks
10 weeks
IV injection

ASO#2
25 mg/kg every week
16 weeks
subcutaneous injection

WT, Gaa$^{-/-}$

Birth   5   5.5   8 (months old)

B Rotarod

Percent change

Month of treatment   0   1   2   3   4
Month of age   5   6   7   8   9

WT
PBS
ASO#2
ASO#2 + ERT
ERT

Gaa$^{-/-}$

****

C

Quadriceps
GYS1
GAPDH
WT PBS ASO#2 ASO#2 + ERT ERT
Gaa$^{-/-}$

Diaphragm
GYS1
GAPDH
WT PBS ASO#2 ASO#2 + ERT ERT
Gaa$^{-/-}$

Heart
GYS1
GAPDH
WT PBS ASO#2 ASO#2 + ERT ERT
Gaa$^{-/-}$

D

Quadriceps

Glycogen (µg/µg protein)

**   ***   ns
        *
            **

WT PBS ASO#2 ASO#2 + ERT ERT
Gaa$^{-/-}$

Heart

Glycogen (µg/µg protein)

ns   *   *

WT PBS ASO#2 ASO#2 + ERT ERT
Gaa$^{-/-}$

E

WT   Gaa$^{-/-}$

PBS   ASO#2

ASO#2 + ERT   ERT

Quadriceps

POMPE DISEASE

PHARMACOLOGICAL INTERVENTION

A Quadriceps

mGys1 mRNA (% PBS) vs Weekly ASO dose (mg/kg)

Gys1 ASO#1
ED_{50} = 45.2 mg/kg

Gys1 ASO#2
ED_{50} = 24.2 mg/kg

B Gastrocnemius

CtrlASO | PBS | 12.5 | 25 | 50 | 100   ASO Dose (mg/kg per week)

GYS1
GAPDH

GYS1:GAPDH

Ctrl ASO | PBS | 12.5 | 25 | 50 | 100
ASO#2

C Quadriceps

PBS | 10 | 12.5 | 25   ASO Dose (mg/kg per week)

GYS1
GAPDH

GYS1:GAPDH

PBS | 10 | 15 | 25
ASO#2

A Cohort 3: 5-month-old adjunct treatment study design

Ctrl ASO

Gys1 ASO#2

Gaa $^{-/-}$