1. Introduction

Glycogenolysis is the major glycogen degradation pathway that involves two cytoplasmic enzymes, glycogen phosphorylase (EC 2.4.1.1) and glycogen debranching enzyme (GDE) [1]. GDE has two independent catalytic activities at different sites on the protein: a 4-alpha-glucosidase (EC 2.4.1.25) activity and an amylo-1, 6-glucosidase (EC 3.2.1.33) activity. Deficiency of GDE in GSD III results in incomplete glycogenolysis and the accumulation of abnormally structured cytoplasmic glycogen in multiple tissues, mainly in liver and muscle. There is a significant clinical variability in GSD III caused by different mutations in the AGL gene. Most patients have both muscle and liver involvement (type IIIa) although some have only liver symptoms (type IIIb) [2,3]. Progressive liver disease with resultant liver fibrosis/cirrhosis and liver failure, myopathy, and cardiomyopathy is a major cause of morbidity and mortality in adult patients. Currently, other than symptomatic management of hypoglycemia and diet interventions, there is no effective treatment for this disease [4].

In addition to the cytosolic glycogenolysis pathway, there is a small percentage of cellular glycogen continually trafficking into the lysosomes and hydrolyzed by the enzyme acid-alpha-glucosidase (GAA, EC 3.2.1.20) [1]. This minor glycogen degradation pathway through lysosomes plays an important role in overall glycogen metabolism because deficiency of GAA causes Pompe disease (GSD II), resulting in an extensive lysosomal glycogen accumulation in skeletal muscle, heart, and the central nerve system [3]. Enzyme replacement therapy (ERT) with recombinant human GAA (rhGAA) is an FDA approved therapy for Pompe disease. We hypothesize that the administration of rhGAA will enhance lysosomal glycogen depletion, facilitate glycogen transport into lysosomes, and ultimately reduce cytoplasmic glycogen accumulation in GSD III. In this study, we tested our hypothesis in a cellular model of GSD III.

2. Materials and methods

Needle skeletal muscle biopsies were obtained from two GSD IIIa patients, a 45 year old male (patient 1, Pt-1) and a 35 year old female (patient 2, Pt-2) at the metabolic clinic of Duke University Medical Center under an institutional review board approved protocol. The diagnosis of GSD IIIa was previously made based on undetectable GDE enzyme activity in their muscle biopsies and two confirmed mutations in the AGL gene at the Duke GSD Laboratory (unpublished data). One-third of each sample was fixed in cold 3% glutaraldehyde for electron microscopy (EM) and high resolution light microscopy.
The remaining tissues were freshly placed in cold DMEM (GIBCO) for the isolation of myoblasts using a standard protocol [6]. The isolated myoblasts were expanded and maintained in high-serum growth medium at 37 °C in a humidified atmosphere of 5% CO2.

Third-passage myoblasts were seeded onto collagen-coated 10-cm dishes in growth medium (Day 1). When cells reached 70–80% confluence (Day 4), differentiation into myotubes (myogenesis) was induced by changing to low-serum differentiation medium [6]. Muscle cells were harvested at different time points (Days 1, 4, 7, 10, 15, and 23) to determine the best treatment time based on glycogen accumulation pattern. A glucose starvation experiment was conducted from Day 13 to Day 15 by removing glucose from medium [7]. For rhGAA treatment, 100 μg of rhGAA (provided by Genzyme) was added to the culture medium (10 ml) on Day 13. After 48 h, cells were washed 3 times with cold phosphate buffered saline (PBS) and then collected with a scraper. GAA activity and glycogen content were determined in the cell lysates as previously described [8].

3. Results

Light microscopic appearance of the patient skeletal muscle biopsies showed abundant, non-membrane-bound glycogen in cytoplasmic pools (Fig. 1A). Under EM, the vast majority of the glycogen was found free in the cytoplasm (not shown) along with a small amount of membrane-bound glycogen (Fig. 1B). Fusion of myoblasts forms large, multinucleated myotubes which mature into myofibers. Long and often branched myotubes were present 3 days (Day 7) after incubation in low-serum differentiation medium. By Day 10, most cells fused into myotubes (not shown). Glycogen content gradually decreased from Day 1 to Day 10, then started to increase and peaked at Day 15 (Fig. 1C). Glucose starvation was conducted from Day 13 to Day 15 and glycogen levels decreased by 28% and 49%, respectively, in the two patient cells, in contrast to an 87% reduction in the normal control cells (Fig. 1D). This indicates incomplete glycogenolysis in the patient cells due to the lack of GDE enzyme activity.

Forty-eight hours after adding rhGAA treatment, GAA activity in medium was decreased from 1905 ± 29 (nmol/h/ml) on Day 13 to 19 ± 1 (nmol/h/ml) on Day 15. In cells, GAA activity was slightly increased for patient 1 but significantly higher for the normal control and patient 2 after rhGAA treatment (Fig. 1E). Interestingly, glycogen content was reduced significantly in both normal (by 35%) and the two patients (by 17% for Pt-1 and 48% for Pt-2, respectively) cells (Fig. 1F).

4. Discussion

Muscle and liver are major affected tissues in human patients with GSD III. Progression of disease can result in patients being wheelchair bound, with end stage liver cirrhosis and severe cardiomyopathy [4].

**Fig. 1.** Analyses of the skeletal muscle biopsies from two GSD IIIa patients, a 45-year-old male (Pt-1) and a 35-year-old female (Pt-2); (A) High-resolution light microscopy demonstrates that purple-staining glycogen is present as non-membrane-bounded cytoplasmic lakes within myocytes by Periodic Acid Schiff (PAS) staining (scale bar = 20 μm). (B) Under EM, occasional lysosomal glycogen (arrow) was also seen in the myocytes (scale bar = 1 μm). (C) Glycogen accumulation pattern revealed that glycogen content was peaked at Day 15 in cultured patient muscle cells. (D) Glucose starvation experiment showed incomplete glycogen utilization in the muscle cells from both GSD IIIa patients compared to a normal control subject (Nor). (E) GAA activity in normal and patient cells 48 h after adding rhGAA treatment. (F) rhGAA significantly reduced glycogen concentration in both normal and patient cells. Mean ± standard deviation is shown in C–F (n = 4). The significance of differences between two different groups was assessed using the two-tailed, equal variance student T-test (*P < 0.001; **P < 0.01; ***P < 0.05).
Dietary interventions include control of hypoglycemia by cornstarch supplements or nocturnal gastric drip feedings and a high protein diet [4]. The benefit of a high protein diet or a combined ketogenic/high protein diet in reversing cardiomyopathy was reported recently as a single case report [9,10]. Thus, there is no definitive therapy for this progressive condition. Challenges with direct ERT and gene therapy for GSD III include the lack of receptor-mediated GDE enzyme uptake and a suitable vector to deliver the large-sized AGL gene.

ERT with rhGAA through mannose-6-phosphate receptor mediated enzyme uptake is an FDA approved therapy for Pompe disease. Although GSD III patients have normal GAA activity in muscle, excessive amounts of enzyme uptake is an FDA approved therapy for Pompe disease. Although rhGAA is a general mechanism but not specific for patients with other GSDs. In future studies, we will test the efficacy of rhGAA treatment in a canine model of GSD IIIa [16].

Conflict of interest

P.S.K. reports receiving research and grant support from Genzyme. B.S., Y.-T.C., and P.S.K. are listed as inventors on a Duke University pending patent application for the use of rhGAA in the treatment of GSD III and other GSDs excluding GSD II. To date, neither Duke University nor the inventors have received any money from rights associated with this pending patent. The other authors declare no conflict of interest.

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References