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# Efficacy of the pharmacologic chaperone migalastat in a subset of male patients with the classic phenotype of Fabry disease and migalastat-amenable variants: data from the phase 3 randomized, multicenter, double-blind clinical trial and extension study

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**Purpose:** Outcomes in patients with Fabry disease receiving migalastat during the phase 3 FACETS trial (NCT00925301) were evaluated by phenotype.

**Methods:** Data were evaluated in two subgroups of patients with migalastat-amenable *GLA* variants: “classic phenotype” ( $n = 14$ ; males with residual peripheral blood mononuclear cell  $\alpha$ -galactosidase A  $<3\%$  normal and multiorgan system involvement) and “other patients” ( $n = 36$ ; males not meeting classic phenotype criteria and all females). Endpoints included estimated glomerular filtration rate (eGFR), left ventricular mass index (LVMI), Gastrointestinal Symptoms Rating Scale diarrhea subscale (GSRS-D), renal peritubular capillary (PTC) globotriaosylceramide (GL-3) inclusions, and plasma globotriaosylsphingosine (lyso-Gb<sub>3</sub>).

**Results:** Baseline measures in the classic phenotype patients suggested a more severe phenotype. At month 24, mean (SD) annualized change in eGFR<sub>CKD-EPI</sub> with migalastat was  $-0.3$  (3.76)

mL/min/1.73 m<sup>2</sup> in the classic phenotype subgroup; changes in LVMI, GSRS-D, and lyso-Gb<sub>3</sub> were  $-16.7$  (18.64) g/m<sup>2</sup>,  $-0.9$  (1.66), and  $-36.8$  (35.78) nmol/L, respectively. At month 6, mean PTC GL-3 inclusions decreased with migalastat ( $-0.8$ ) and increased with placebo (0.3); switching from placebo to migalastat, PTC inclusions decreased by  $-0.7$ . Numerically smaller changes in these endpoints were observed in the other patients.

**Conclusion:** Migalastat provided clinical benefit to patients with Fabry disease and amenable variants, regardless of disease severity.

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## INTRODUCTION

Fabry disease (OMIM 301500) is a devastating, rare, and progressive X-linked lysosomal disorder caused by pathogenic variants in the *GLA* gene, resulting in functional deficiency of  $\alpha$ -galactosidase A ( $\alpha$ -Gal A).<sup>1,2</sup> This causes accumulation of globotriaosylceramide (GL-3 or Gb<sub>3</sub>) within lysosomes in various tissues, producing a wide variety of clinical symptoms and manifestations, including peripheral neuropathy, cardiovascular disease, stroke, end-stage renal disease, gastrointestinal

disorders, and premature death.<sup>1,2</sup> Although some *GLA* variants do not appear to cause disease,<sup>1</sup> more than 1000 disease-causing variants have been reported, of which ~60% are missense.<sup>3,4</sup> In males, disease severity is generally inversely related to residual  $\alpha$ -Gal A activity with variants resulting in near or complete loss of function associated with the “classic” early-onset form of the disease.<sup>1</sup> Females generally have later symptom onset than males, but can develop the classic phenotype due to skewed X chromosome inactivation pattern.<sup>5,6</sup>

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Enzyme replacement therapy (ERT) has been the current standard treatment for Fabry disease. ERT has been shown to reduce disease substrate (i.e., GL-3 and plasma globotriaosylsphingosine [lyso-Gb<sub>3</sub>]) and symptoms.<sup>1,2,7–9</sup> However, response to treatment is dependent on several factors including disease severity, and the amount of organ damage present at treatment initiation.<sup>10–13</sup> Furthermore, lifelong, biweekly intravenous ERT infusions are a burden, which can result in delayed treatment initiation and reduced compliance. Infusion reactions to ERT have been reported,<sup>10</sup> and development of anti-drug antibodies can reduce the efficacy of ERT, as demonstrated in a 5-year retrospective analysis in which 40% of males had serum-mediated antibody inhibition of agalsidase activity.<sup>14</sup>

Migalastat, a first-in-class, orally administered small molecule, is a pharmacological chaperone that binds to and stabilizes *amenable* mutant forms of  $\alpha$ -Gal A, facilitating lysosomal trafficking and increasing lysosomal enzyme activity.<sup>4,15–19</sup> Amenable mutant forms of  $\alpha$ -Gal A are identified using the migalastat amenability assay, which measures migalastat-induced changes in human embryonic kidney (HEK) cells that are transfected with DNA plasmids containing *GLA* variants.<sup>4</sup> Established criteria for amenability are an increase in  $\alpha$ -Gal A activity  $\geq 1.2$ -fold above baseline and an absolute increase of  $\geq 3\%$  of wild-type (WT)  $\alpha$ -Gal A after incubation with 10  $\mu$ M migalastat.<sup>4</sup>

In patients with amenable variants, migalastat is a treatment option. In the phase 3, placebo-controlled FACETS study (ClinicalTrials.gov; NCT00925301) in patients with Fabry disease and amenable variants who were either ERT-naïve or had not received ERT within the past 6 months, migalastat led to reduced substrates in kidney and plasma, stabilized renal function, reduced cardiac mass, and improved gastrointestinal symptoms.<sup>20</sup> In the phase 3, active-controlled ATTRACT study (ClinicalTrials.gov; NCT01218659) in ERT-experienced patients, migalastat was associated with a reduction in cardiac mass, had similar effects on renal function compared with ERT, and was generally well-tolerated.<sup>19</sup> These results led to the approval of migalastat in the European Union, Switzerland, Australia, Republic of Korea, Israel, and Japan for the treatment of Fabry disease in patients over 16 years of age, with amenable *GLA* variants and GFR  $\geq 30$  mL/min/1.73 m<sup>2</sup> (refs. <sup>15,21</sup>). In addition, migalastat is approved in Canada and the United States for the long-term treatment of Fabry disease in adults ( $\geq 18$  years old) with amenable *GLA* variants.<sup>22,23</sup>

The objective of the present analyses was to assess the clinical benefit of migalastat in the subset of male patients with the classic phenotype in the FACETS trial (i.e., multiorgan system involvement and residual peripheral blood mononuclear cell [PBMC]  $\alpha$ -Gal A activity  $< 3\%$  of normal).<sup>20</sup> Results in male patients not meeting classic phenotype criteria and all female patients were also assessed.

## MATERIALS AND METHODS

### In vitro assays

Methodology for the Good Laboratory Practice (GLP)-validated in vitro assay in HEK-293 cells has been published.<sup>4</sup>

In brief, plasmids containing WT or mutated  $\alpha$ -Gal A complementary DNA (cDNA) were used to transfect HEK-293 cells; transfected cells were then incubated in the presence or absence of 10  $\mu$ mol/L migalastat for 5 days, after which cell lysates were assayed for  $\alpha$ -Gal A activity. Variants meeting the prespecified criteria for amenability in the assay were categorized as amenable.<sup>4</sup>

### FACETS phase 3 study design

The multicenter, phase 3, randomized, placebo-controlled, double-blind FACETS trial to evaluate the efficacy and safety of migalastat in patients with Fabry disease and amenable variants has been described in detail.<sup>20</sup> Briefly, stage 1 of the study consisted of a 6-month, double-blind treatment period in which patients were randomly assigned to receive migalastat 150 mg or placebo every other day. After stage 1, patients could receive open-label migalastat 150 mg every other day for an additional 6 months, and another 12 months during an open-label extension.

The study was approved by the institutional review board or ethics committee at each participating center and was conducted in accordance with the International Conference on Harmonization and Good Clinical Practice guidelines and the principles of the Declaration of Helsinki; all patients provided written informed consent.

### Clinical analyses

Two subgroups of patients with amenable variants enrolled in the FACETS study were identified for evaluation: “classic phenotype” and “other patients.” The classic phenotype subgroup included male patients with the classic phenotype of Fabry disease and migalastat-amenable variants; these patients had residual PBMC  $\alpha$ -Gal A activity  $< 3\%$  normal and multiorgan system involvement, defined as two or more of the following organ system involvements: renal, cardiac, central nervous system, neuropathic pain, and gastrointestinal symptoms.<sup>20</sup> The other patients subgroup included male patients with migalastat-amenable variants who did not meet the criteria defined above and all females.<sup>17,20</sup>

Changes in measurements of disease severity from baseline are reported. Detailed methodology has been published.<sup>20</sup> Annualized rates of change from baseline to month 24 in estimated glomerular filtration rate using the Chronic Kidney Disease Epidemiology Collaboration equation (eGFR<sub>CKD-EPI</sub>) and GFR measured using iohexol clearance (mGFR<sub>iohexol</sub>) were calculated based on the slope of the linear regression between the observed values and the assessment times. Mean change from baseline to month 24 in left ventricular mass index (LVMI) was calculated based on 2D or M-mode echocardiograms assessed by a single, blinded reader at a central laboratory. Changes from baseline in number of GL-3 inclusions per renal peritubular capillary (PTC) were based on kidney biopsies performed at baseline, month 6, and month 12, which were assessed per BLISS methodology performed by 3 independent, blinded pathologists.<sup>20,24</sup> The same three pathologists qualitatively assessed GL-3 changes in podocytes,

endothelial cells, and mesangial cells. Change from baseline to month 24 in plasma lyso-Gb<sub>3</sub> levels was based on measurements of retained iohexol plasma samples from each time point. Twenty-four-hour urine protein levels were measured by a central laboratory. PBMC  $\alpha$ -Gal A activity was measured with a fluorescence assay of PBMC lysates and normalized to total protein.<sup>4</sup>

The Gastrointestinal Symptoms Rating Scale was used to assess change from baseline to month 24 in diarrhea symptoms (GSRS-D). A response in the GSRS-D was defined as a reduction  $\geq 0.33$ . The minimal clinically important difference (MCID) of 0.33 was based on estimates from the literature on several non-Fabry gastrointestinal disorders in which diarrhea is prominent.<sup>25,26</sup> Distribution-based estimates of MCID in Fabry disease were based on the change from baseline in the placebo arm of the FACETS study, generating an MCID of 0.35 based on half the standard deviation,<sup>20</sup> supporting an MCID for GSRS-D of 0.33 in Fabry patients.

For all outcomes except PTC GL-3 inclusions, data are pooled for patients within the designated subgroups (classic phenotype and other patients) regardless of treatment allocation for the first 6 months (migalastat or placebo). Change from baseline to month 24 data are the change from baseline to month 24 for patients randomized to migalastat in stage 1 of the study and the change from month 6 (start of migalastat) to month 24 in patients randomized to placebo in stage 1.

All data were derived from post hoc analyses and formal statistical testing was not performed. Data were analyzed descriptively using mean, standard deviation (SD) or standard error (SE), and 95% confidence intervals (CIs).

## RESULTS

### In vitro activity

For the *GLA* variants found in male patients enrolled in this study who met the criteria for the classic phenotype, the mean absolute increase in  $\alpha$ -Gal A activity with 10  $\mu$ M migalastat in vitro was 24.8% of WT activity; mean activity increased by 8.2-fold above baseline (Table 1). For all other variants, mean absolute increase in  $\alpha$ -Gal A activity in vitro was 24.5% of WT; mean activity increased by 6.8-fold above baseline (Table 1).

### Clinical data

#### Baseline characteristics

The classic phenotype subgroup includes 14 males with the classic phenotype (16–62 years old). The other patients subgroup includes 36 patients (4 males with nonclassic phenotype [33–58 years old], 32 females [27–68 years old]). Disease severity of patients in the classic phenotype subgroup was evident from baseline measurements of variables and medical history, indicating multisystem involvement (Table 2 and Fig. 1a–e). Mean (SD) baseline eGFR<sub>CKD-EPI</sub> values were 87.8 (33.6) and 95.3 (19.6) mL/min/1.73 m<sup>2</sup>, and mean (SD) baseline mGFR<sub>iohexol</sub> values were 78.6 (22.9) and 88.2 (22.0)

mL/min/1.73 m<sup>2</sup> in the classic phenotype and other patients subgroup, respectively.

*GLA* variants in male patients with the classic phenotype were p.Ile253Thr ( $n = 2$ ), p.Pro259Arg ( $n = 2$ ), p.Gly183Asp, p.Leu243Phe, p.Cys174Arg, p.Asp55Val/Gln57Leu, p.Gly144-Val, p.Arg301Gln, p.Gly373Ser, p.Asp322Glu, p.Gly325Arg, and p.Tyr216Cys ( $n = 1$  each).

Six (43%) male patients with the classic phenotype were taking angiotensin-converting enzyme inhibitors, angiotensin receptor blockers, or renin inhibitors at baseline; the dosages of these medications had been stable for  $\geq 4$  weeks before the screening visit and did not change during the study.

### Renal function

The mean (SD) annualized rate of change in eGFR<sub>CKD-EPI</sub> from baseline to month 24 was  $-0.3$  (3.76) mL/min/1.73 m<sup>2</sup> (95% CI:  $-2.80$ , 2.25;  $n = 11$ ) in the classic phenotype subgroup and  $-0.3$  (4.47) mL/min/1.73 m<sup>2</sup> (95% CI:  $-2.0$ , 1.4;  $n = 30$ ) in other patients (Fig. 1a; individual values for male patients with the classic phenotype in Table S1). Median annualized rates of change in the classic phenotype and other patients subgroups were 0.25 mL/min/1.73 m<sup>2</sup> (min, max:  $-8.6$ , 4.3) and 0.33 mL/min/1.73 m<sup>2</sup> (min, max:  $-13.8$ , 7.4), respectively. Mean (SD) annualized rate of change in mGFR<sub>iohexol</sub> was  $-3.0$  (6.04) mL/min/1.73 m<sup>2</sup> (95% CI:  $-7.65$ , 1.64;  $n = 9$ ) in the classic phenotype subgroup and  $-1.0$  (8.66) mL/min/1.73 m<sup>2</sup> (95% CI:  $-4.39$ , 2.33;  $n = 28$ ) in other patients. One male patient with the classic phenotype had a reduction of  $-16$  mL/min/1.73 m<sup>2</sup> from month 6 to month 24. Median (range) change in mGFR<sub>iohexol</sub> from baseline to month 24 was  $-1.03$  mL/min/1.73 m<sup>2</sup> (min, max:  $-16.2$ , 3.0) in the classic phenotype subgroup and  $-0.34$  mL/min/1.73 m<sup>2</sup> (min, max:  $-20.4$ , 15.8) in the other patients subgroup.

Twenty-four-hour urine protein levels showed small increases from baseline to month 24 in both subgroups. Baseline mean (SD) 24-hour urinary protein excretion values were 767.7 (710.7) (95% CI: 357.4, 1178.0;  $n = 14$ ) and 360.6 (508.7) (95% CI: 183.1, 538.1;  $n = 34$ ) mg/24 hours in the classic phenotype and other patients subgroups, respectively. Mean (SD) changes from baseline to month 24 in 24-hour urine protein were 271.3 (419.6) (95% CI:  $-79.5$ , 622.0;  $n = 8$ ) and 168.0 (494.4) (95% CI:  $-20.0$ , 356.1;  $n = 29$ ) mg/24 hours in the classic phenotype and other patients subgroups, respectively.

Results were similar to those reported for the overall amenable FACETS population, in which migalastat stabilized eGFR in patients with amenable *GLA* variants, with mean (SE) annualized changes from baseline to month 24 in eGFR<sub>CKD-EPI</sub> and mGFR<sub>iohexol</sub> of  $-0.3$  (0.66) and  $-1.5$  (1.33) mL/min/1.73 m<sup>2</sup>, respectively.<sup>20</sup>

### Cardiac geometry

Migalastat led to reductions in LVMI in both subgroups. Of note, mean (SD) baseline LVMI values were 114.3 (27.3) g/m<sup>2</sup> in male patients with the classic phenotype ( $n = 14$ ) and 88.2 (32.3) g/m<sup>2</sup> in other patients ( $n = 30$ ). Seven (50%) males with the classic phenotype and 4 (11%) in the other patients

**Table 1** Effect of migalastat on  $\alpha$ -Gal A activity by GLA variant<sup>4</sup>  
 Measured in PBMC at baseline (month 0)<sup>a</sup> Measured in HEK-293 cell lysate

$\alpha$ -Gal A mutant form	Nucleotide change	-Migalastat		+Migalastat		Absolute increase at 10 $\mu$ M (%WT)	$\alpha$ -Gal A activity at 10 $\mu$ M (fold over baseline)		
Amino acid change		$\alpha$ -Gal A activity (nmol/mg/h)	% WT	$\alpha$ -Gal A activity (nmol/mg/h)	% WT				
<b>GLA variants in males meeting criteria for the classic phenotype<sup>b</sup></b>									
p.Asp55Val/ Gln57Leu	c.164 A>T; c.170 A>T	BLQ	0.2	9913	N/A	2526 $\pm$ 287 <sup>c</sup>	8.0 $\pm$ 0.5	8.0	NC
p.Gly144Val	c.431 G>T	0.05	0.2	253 $\pm$ 21	0.8 $\pm$ 0.1	2924 $\pm$ 335 <sup>c</sup>	9.2 $\pm$ 1.2	8.4	11.58
p.Cys174Arg	c.520 T>C	0.28	1.3	4671 $\pm$ 270	14.3 $\pm$ 0.6	16,505 $\pm$ 393 <sup>c</sup>	51.6 $\pm$ 2.0	37.4	3.53
p.Gly183Asp	c.548 G>A	0.04	0.0	207 $\pm$ 15	0.7 $\pm$ 0.1	6074 $\pm$ 212 <sup>c</sup>	19.1 $\pm$ 1.0	18.4	29.31
p.Tyr216Cys	c.647 A>G	BLQ	0.0	673 $\pm$ 38	2.0 $\pm$ 0.1	7003 $\pm$ 305 <sup>c</sup>	20.7 $\pm$ 0.8	18.7	10.40
p.Leu243Phe	c.729 G>C	0.19	1.1	2694 $\pm$ 117	7.9 $\pm$ 0.3	14,370 $\pm$ 618 <sup>c</sup>	42.3 $\pm$ 1.4	34.4	5.33
p.Ile253Thr <sup>c</sup>	c.758 T>C	0.63/0.58	2.9/2.6	11,287 $\pm$ 506	38.9 $\pm$ 3.0	23,417 $\pm$ 1077 <sup>c</sup>	80.2 $\pm$ 5.9	41.3	2.07
p.Pro259Arg <sup>c</sup>	c.776 C>G	0.44/0.6	3.3/3.1	6681 $\pm$ 364	23.3 $\pm$ 2.3	17,645 $\pm$ 515 <sup>c</sup>	60.3 $\pm$ 3.8	37.0	2.64
p.Arg301Gln	c.902 G>A	0.42	2.5	1914 $\pm$ 52	5.5 $\pm$ 0.2	15,547 $\pm$ 353 <sup>c</sup>	44.5 $\pm$ 1.0	39.0	8.12
p.Asp322Glu	c.966 C>A	0.2	0.9	2398 $\pm$ 141	6.7 $\pm$ 0.2	9554 $\pm$ 667 <sup>c</sup>	26.8 $\pm$ 1.3	20.0	3.98
p.Gly325Arg	c.973 G>C	0.07	0.3	909 $\pm$ 31	2.6 $\pm$ 0.1	9244 $\pm$ 417 <sup>c</sup>	26.6 $\pm$ 1.4	24.0	10.17
p.Gly373Ser	c.1117 G>A	0.29	0.8	1544 $\pm$ 69	4.8 $\pm$ 0.3	5128 $\pm$ 288 <sup>c</sup>	15.7 $\pm$ 0.8	10.9	3.32
<b>GLA variants in males that did not meet criteria for the classic phenotype</b>									
p.Asp33Gly	c.98 A>G	1.42	6.5	9913 $\pm$ 600	29.3 $\pm$ 1.8	24,033 $\pm$ 865 <sup>c</sup>	70.6 $\pm$ 2.4	41.3	2.42
p.Ala156Thr <sup>d</sup>	c.466 G>A	NA	NA	907 $\pm$ 31	2.8 $\pm$ 0.1	7034 $\pm$ 289 <sup>c</sup>	21.9 $\pm$ 0.9	19.1	7.75
p.Asp244Asn <sup>e</sup>	c.730 G>A	NA	NA	10,317 $\pm$ 386	30.9 $\pm$ 1.6	16,321 $\pm$ 402 <sup>c</sup>	48.7 $\pm$ 1.7	17.8	1.58
p.Arg356Trp	c.1066 C>T	1.22	1.7	3526 $\pm$ 240	11.0 $\pm$ 0.7	15,570 $\pm$ 830 <sup>c</sup>	49.1 $\pm$ 2.6	38.1	4.42
<b>GLA variant in female patients</b>									
p.Leu36Trp	c.107 T>G	—	—	241 $\pm$ 19	0.7 $\pm$ 0.1	5182 $\pm$ 463 <sup>c</sup>	16.6 $\pm$ 2.1	15.9	21.49
p.Gly85Asp	c.254 G>A	—	—	888 $\pm$ 38	2.7 $\pm$ 0.1	4534 $\pm$ 135 <sup>c</sup>	14.3 $\pm$ 0.9	11.6	5.10
p.Arg112His	c.335 G>A	—	—	845 $\pm$ 39	2.6 $\pm$ 0.1	5583 $\pm$ 215 <sup>c</sup>	17.4 $\pm$ 0.8	14.8	6.61
p.Met187Ile	c.561 G>A	—	—	1775 $\pm$ 57	5.1 $\pm$ 0.2	10,824 $\pm$ 555 <sup>c</sup>	30.7 $\pm$ 1.1	25.6	6.10
p.Pro205Thr	c.613 C>A	—	—	4802 $\pm$ 230	14.4 $\pm$ 0.9	16,371 $\pm$ 647 <sup>c</sup>	48.8 $\pm$ 2.2	34.4	3.41
p.Gly258Arg	c.772 G>C	—	—	9558 $\pm$ 348	32.6 $\pm$ 2.1	22,630 $\pm$ 801 <sup>c</sup>	78.1 $\pm$ 5.8	45.5	2.37
p.Leu300Pro	c.899 T>C	—	—	1277 $\pm$ 38	3.7 $\pm$ 0.1	13,219 $\pm$ 412 <sup>c</sup>	37.9 $\pm$ 1.2	34.2	10.35
p.Pro293Thr	c.877 C>A	—	—	229 $\pm$ 20	0.7 $\pm$ 0.1	4488 $\pm$ 327 <sup>c</sup>	13.3 $\pm$ 1.1	12.7	19.60
p.Phe 295Cys	c.884 T>G	—	—	1196 $\pm$ 38	3.4 $\pm$ 0.1	5051 $\pm$ 190 <sup>c</sup>	14.5 $\pm$ 0.6	11.1	4.22
p.Gly271Ser; Asp313Tyr <sup>f</sup>	c.811 G>A; c.937 G>T	—	—	BLQ	N/A	877 $\pm$ 18 <sup>c</sup>	3.0 $\pm$ 0.2	3.0	NC
p.Ile317Thr	c.950 T>C	—	—	2298 $\pm$ 338	6.5 $\pm$ 0.6	7812 $\pm$ 530 <sup>c</sup>	23.6 $\pm$ 1.0	17.0	3.40
p.Asp264Tyr	c.790 G>T	—	—	143 $\pm$ 13	0.5 $\pm$ 0.0	1842 $\pm$ 100 <sup>c</sup>	6.2 $\pm$ 0.3	5.7	12.89
p.Gly260Ala	c.779 G G>C	—	—	2221 $\pm$ 142	7.5 $\pm$ 0.6	10,749 $\pm$ 403 <sup>c</sup>	37.4 $\pm$ 3.1	29.9	4.84
p.Met284Thr	c.851 T>C	—	—	606 $\pm$ 40	1.7 $\pm$ 0.1	5050 $\pm$ 268 <sup>c</sup>	14.3 $\pm$ 0.6	12.6	8.33
p.Gly183Asp	c.548 G>A	—	—	207 $\pm$ 15	0.7 $\pm$ 0.1	6074 $\pm$ 212 <sup>c</sup>	19.1 $\pm$ 1.0	18.4	29.31

**Table 1** continued

α-Gal A mutant form	Nucleotide change	Measured in PBMC at baseline (month 0) <sup>a</sup>				Measured in HEK-293 cell lysate				α-Gal A activity at 10 μM (fold over baseline)
		-Migalastat		% WT		-Migalastat		% WT		
		α-Gal A activity (nmol/mg/h)	% WT	α-Gal A activity (nmol/mg/h)	% WT	α-Gal A activity (nmol/mg/h)	% WT	α-Gal A activity (nmol/mg/h)	% WT	
p.Arg301Gln	c.902 G>A	—	—	1914 ± 52	5.5 ± 0.2	15,547 ± 353 <sup>c</sup>	44.5 ± 1.0	39.0	8.12	
p.Ile253Thr	c.758 T>C	—	—	11,287 ± 506	38.9 ± 3.0	23,417 ± 1077 <sup>c</sup>	80.2 ± 5.9	41.3	2.07	
p.Asp322Glu	c.966 C>A	—	—	2398 ± 141	6.7 ± 0.2	9554 ± 667 <sup>c</sup>	26.8 ± 1.3	20.0	3.98	
p.Ile270Thr	c.809 T>C	—	—	1846 ± 119	6.3 ± 0.5	12,416 ± 377 <sup>c</sup>	42.8 ± 3.0	36.5	6.73	
p.Gly325Arg	c.973 G>C	—	—	909 ± 31	2.6 ± 0.1	9244 ± 417 <sup>c</sup>	26.6 ± 1.4	24.0	10.17	
p.Tyr216Cys	c.647 A>G	—	—	673 ± 38	2.0 ± 0.1	7003 ± 305 <sup>c</sup>	20.7 ± 0.8	18.7	10.40	
p.Pro259Arg	c.776 C>G	—	—	6681 ± 364	23.3 ± 2.3	17,645 ± 515 <sup>c</sup>	60.3 ± 3.8	37.0	2.64	
p.Met284Thr	c.851 T>C	—	—	606 ± 40	1.7 ± 0.1	5050 ± 268 <sup>c</sup>	14.3 ± 0.6	12.6	8.33	
p.Gly258Arg	c.772 G>C	—	—	9558 ± 348	32.6 ± 2.1	22,630 ± 801 <sup>c</sup>	78.1 ± 5.8	45.5	2.37	

α-Gal A α-galactosidase A, BLQ below the limit of quantification (<142 nmol/mg/h), NA not assessed, PBMC peripheral blood mononuclear cells, WT wild-type. This table lists the unique GLA variants occurring in the different patient subgroups, with some variants found in ≥1 patient; therefore, the numbers of variants do not necessarily reflect patient numbers. These data have been previously published by Benjamin et al.<sup>4</sup> Data on the variant relevant to the current paper have been collated from Table 15 (HEK-293) and Table 45 (PBMC) of the Supplementary Material from Benjamin et al. Data are expressed as the mean ± SEM of 20 data points: mutant α-Gal A activity is expressed as a percentage of the α-Gal A activity measured in WT cell lysates incubated without migalastat (-Migalastat) assayed in parallel. Absolute increase at 10 μM (%WT) = the percentage of wild-type α-Gal A activity with 10 μM migalastat (+Migalastat) minus the baseline (-Migalastat) percentage of wild-type α-Gal A activity. BLQ indicates that the mean α-Gal A activity (mean of  $n = 20$ ) was below the limit of quantification (nmol/mg/h; value is equal to 3<sup>c</sup> standard deviation of the pDNA activity in nmol/mg/h after vector subtraction across 128 samples assayed in ten method validation experiments).

α-Gal A activity at 10 μM (fold over baseline) = α-Gal A activity in mutant-transfected cell lysate with 10 μM migalastat / α-Gal A activity in mutant-transfected cell lysate without migalastat. Statistically significant differences in α-Gal A activity without migalastat ( $n = 20$ ) versus with 10 μM migalastat ( $n = 20$ ) were determined using a one-tailed Mann-Whitney U nonparametric test.

<sup>a</sup>PBMC data are only shown for male patients.

<sup>b</sup>Criteria for the classic phenotype: male, PBMC α-Gal A activity <3% of normal, multiorgan disease.

<sup>c</sup>α-Gal A mutant forms that show a statistically significant increase in α-Gal A activity after incubation with 10 μM migalastat.  $P < 0.001$ .

<sup>d</sup>Two male patients with the classic phenotype included in the FACETS trial had this variant.

<sup>e</sup>The PBMC α-Gal A value at baseline from one male patient (p.A156T) was implausible because it was higher than any baseline value from all other male patients in the study (in fact, the baseline value is within the typical range of baseline values from female patients in AT1001-01), and it was higher than any one of this male's other visits. Therefore, this patient was excluded from this analysis and is not represented in this table.

<sup>f</sup>The PBMC α-Gal A value at baseline was missing for one male patient (p.D244N) treated with migalastat in stage 1 and stage 2. Therefore, this patient was excluded from this analysis and is not represented in this table.

<sup>g</sup>Variant not reported as Fabry disease-associated.



**Table 2** Baseline characteristics of male patients with the classic phenotype  
Male patients with the classic phenotype

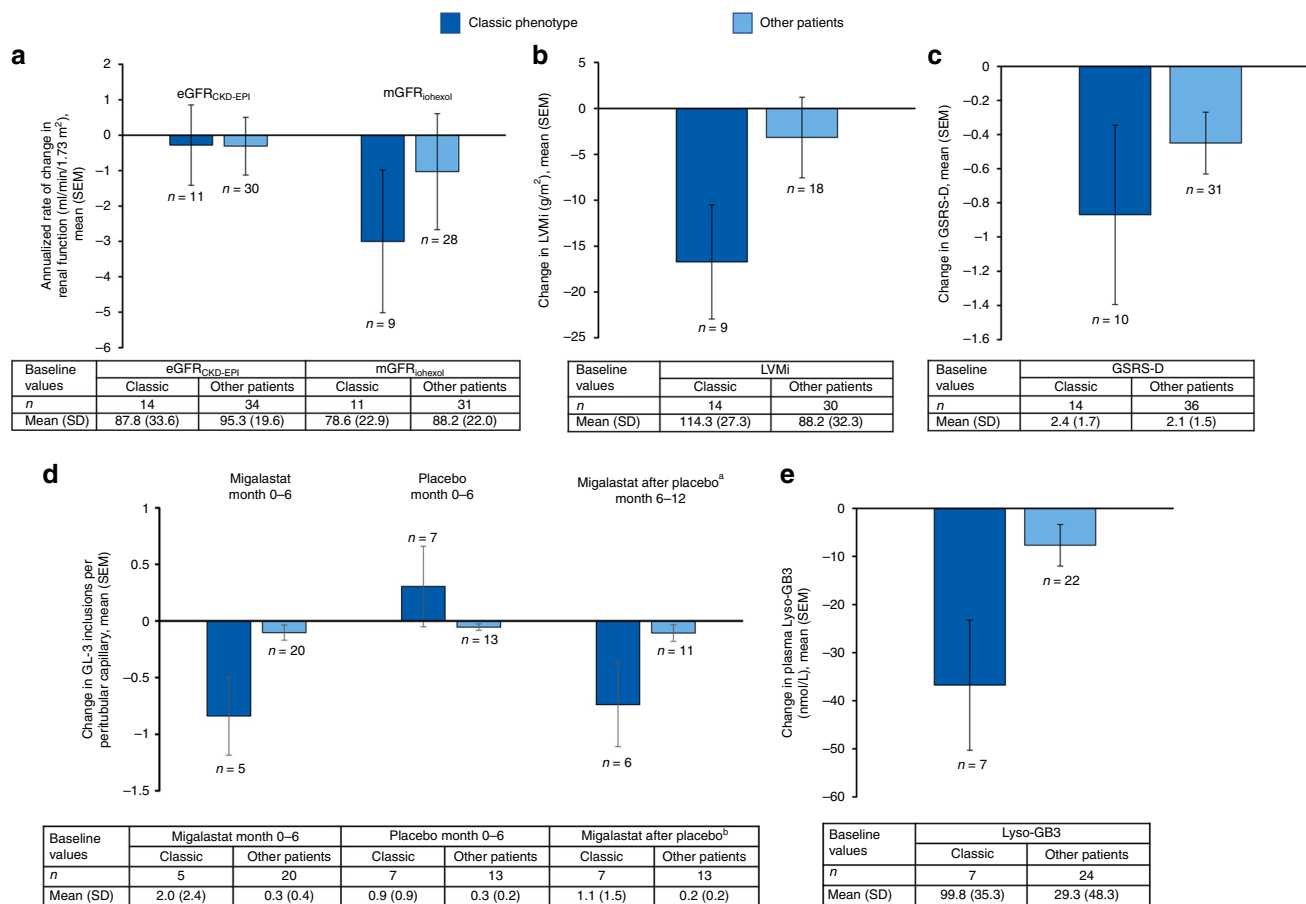
	1	2	3	4	5	6	7	8	9	10	11	12	13	14
Variant (amino acid change)	p. I253T	p. I253T	p. C174R	p. G144V	p. D322E	p. G325R	p. Y216C	p. G183D	p. L243F	p. p.D55V; p.Q57L	p. R301Q	p. G373S	p. P259R	p. P259R
<b>Baseline information</b>														
Age (years)	62	61	27	35	60	16	34	35	45	25	56	48	45	45
Weight (kg)	74	58	92.7	66.5	97.07	64	90.5	70	67	100.6	77.6	111.2	101.8	62.7
Time since Fabry diagnosis (years)	4.6	0.3	2.6	13.4	2.4	0.4	6.2	23.9	9.1	2.6	0.6	0.5	9.9	8.9
Previous use of ERT	Yes	No	No	No	No	No	Yes	Yes	Yes	No	No	No	Yes	Yes
Blood pressure (SBP/DBP, mm Hg)	120/80	NA	120/70	110/63	136/84	130/70	130/90	125/62	117/63	137/96	122/94	119/66	140/80	110/70
Urine protein (mg/24 hours)	331	1900	1584	240	918	242	400	150	161	198	2351	602	335	1909
eGFR <sub>CKD-EPI</sub> (mL/min/1.73 m <sup>2</sup> )	86.78	53.79	44.12	104.9	41.36	151.35	119.22	109.68	102.24	98.08	55.71	64.55	105.36	86.34
LVMi (g/m <sup>2</sup> )	176.18	142.78	77.36	114.93	140.79	77.86	102.06	101.77	105.02	83.42	137.47	121.68	115.86	105.77
Urine GL-3 (ng/mg creatinine)	1975	4063.6	1090	2176.2	4163.6	2119.2	3418.9	641.4	4196.6	750.9	1002.4	847.6	2548.6	5245.9
Plasma lyso-Gb3 (nmol/L)	NA	NA	NA	119.7	81.7	NA	128	NA	109	92.2	NA	NA	113.3	53.3
GSRS-D	1	1	1	4	2	4.7	1.3	1	1	2.7	1	3.3	4	6
WBC α-Gal A activity (4 MU/h/mg)	0.63	0.58	0.28	0.05	0.2	0.07	BLQ	0.04	0.19	BLQ	0.42	0.29	0.6	0.44
Use of ACEI/ARB/RI	Yes	No	Yes	No	No	No	No	Yes	Yes	No	Yes	Yes	No	No
<b>Medical history<sup>a</sup></b>														
Acroparesthesia	No	No	No	Yes	No	No	No	Yes	Yes	No	No	No	No	No
Angiokeratoma	No	No	No	No	Yes	No	Yes	Yes	Yes	Yes	No	Yes	Yes	Yes
Cornea verticillata	No	No	No	No	No	No	No	No	Yes	No	No	No	No	No
Cardiac event <sup>b</sup>	Yes	No	Yes	Yes	Yes	No	No	Yes	Yes	Yes	Yes	Yes	Yes	Yes
TIA/stroke event	Yes	No	No	No	Yes	No	No	No	No	No	No	No	No	Yes
Hearing loss	Yes	No	No	Yes	Yes	No	No	Yes	Yes	No	No	No	No	Yes
Hypertension	No	No	Yes	No	Yes	No	No	No	No	No	No	Yes	No	No
Hypohidrosis	No	No	No	Yes	No	Yes	Yes	Yes	No	Yes	No	No	No	No
GI symptoms <sup>c</sup>	No	No	Yes	Yes	Yes	Yes	Yes	No	No	Yes	Yes	Yes	No	Yes

ACEI angiotensin-converting-enzyme inhibitor, ARB angiotensin II receptor blocker, BLQ below the limit of quantification, DBP diastolic blood pressure, eGFR<sub>CKD-EPI</sub> estimated glomerular filtration rate using the Chronic Kidney Disease Epidemiology Collaboration equation, ERT enzyme replacement therapy, GI gastrointestinal, GSRS-D Gastrointestinal Symptom Rating Scale diarrhea domain, LVMi left ventricular mass index, NA not assessed, RI renin inhibitor, SBP systolic blood pressure, TIA transient ischemic attack, WBC white blood cell.

<sup>a</sup>“Yes” indicates reported and “No” indicates not reported. However, medical history may not have been completely reported; therefore, “No” does not necessarily indicate absence of the complication.

<sup>b</sup>Cardiac events include left ventricular hypertrophy, heart failure, heart valve regurgitation, myocardial infarction, bradycardia, right bundle branch block, atrial fibrillation, and tachycardia.

<sup>c</sup>GI symptoms include diarrhea, nausea, vomiting, gastroesophageal reflux disease, and abdominal pain/distension.



**Fig. 1** Measurements of disease severity and treatment effect in the classic phenotype and other patient subgroups. (a) Mean annualized rate of change in estimated glomerular filtration rate (eGFR) and measured glomerular filtration rate (mGFR) from baseline to month 24. (b) Mean change in LVMI from baseline to month 24. (c) Mean change in GRS-D from baseline to month 24. (d) Mean change from baseline to month 12 in average number of GL-3 inclusions per peritubular capillary. Within each subgroup, patients are grouped according to treatment allocation (migalastat to migalastat or placebo to migalastat). (e) Mean change from baseline to month 24 in plasma lyso-Gb<sub>3</sub>. <sup>a</sup>Group is comprised of patients who switched from placebo to migalastat at month 6. <sup>b</sup>Data are mean (SD) number of GL-3 inclusions per peritubular capillary at month 6 (i.e., after 6 months of placebo treatment). The upper cutoff value for normal plasma lyso-Gb<sub>3</sub> level is 2.4 nmol/L.<sup>38</sup> eGFR<sub>CKD-EPI</sub> estimated glomerular filtration rate using the Chronic Kidney Disease Epidemiology Collaboration equation, GL-3 globotriaosylceramide, GRS-D Gastrointestinal Symptoms Rating Scale diarrhea subscale, LVMI left ventricular mass index, lyso-Gb<sub>3</sub> plasma globotriaosylsphingosine, mGFR<sub>iohexol</sub> glomerular filtration rate measured using iohexol clearance.

subgroup had left ventricular hypertrophy (LVH; >115 g/m<sup>2</sup> for males; >95 g/m<sup>2</sup> for females) at baseline. Mean (SD) change from baseline to month 24 in LVMI was −16.7 (18.64) g/m<sup>2</sup> (95% CI: −31.1, −2.4; n = 9) in males with the classic phenotype and −3.2 (18.66) g/m<sup>2</sup> (95% CI: −12.5, 6.1; n = 18) in other patients (Fig. 1b; individual values for male patients with the classic phenotype in Table S1).

By contrast, LVMI change in the overall migalastat-treated amenable population in FACETS was −7.7 g/m<sup>2</sup> (95% CI: −15.4, −0.01; n = 27) after 18–24 months of treatment with migalastat, with larger decreases in the subgroup of patients with LVH (−18.6 g/m<sup>2</sup> [95% CI: −38.2, 1.0]; −11.5%; n = 8) (ref. 20).

#### Gastrointestinal signs and symptoms

Mean (SD) GRS-D scores at baseline were 2.4 (1.69; n = 14) in the classic phenotype subgroup and 2.1 (1.49; n = 36) in the other patients subgroup. Eight (57%) male patients with

the classic phenotype had diarrhea symptoms at baseline (GRS-D score ≥1). In the classic phenotype subgroup, 6 months of treatment (stage 1) with migalastat improved (reduced) scores in GRS-D (mean [SD] change from baseline −0.3 [0.77]; 95% CI: −1.0, 0.4; n = 7), whereas patients treated with placebo had a small increase (0.2 [0.46]; 95% CI: −0.2, 0.7; n = 7).

Male patients with the classic phenotype benefited from continuing treatment with migalastat or by switching from placebo to migalastat. By 24 months, the mean (SD) change from baseline in diarrhea symptoms was −0.9 (1.66; n = 10) (Fig. 1c; individual values in Table S1). Seven of the 8 (88%) males with the classic phenotype that had diarrhea symptoms at baseline achieved an MCID reduction of 0.33 in GRS-D. The corresponding change in other patients was −0.5 (1.01; n = 31) at month 24.

These results are similar to those of the overall amenable FACETS population, in which patients with amenable GLA

variants treated with migalastat had mean changes in GSRS-D of  $-0.3$  (versus  $0.2$  in the placebo group) at month 6 and  $-0.5$  (95% CI:  $-0.9, -0.1$ ) at month 24 (ref.<sup>20</sup>).

#### PTC GL-3 inclusions

During the first 6 months of the study, the mean (SD) number of GL-3 inclusions per PTC decreased from baseline in males with the classic phenotype treated with migalastat ( $-0.8$  [ $-0.78$ ]; 95% CI:  $-1.8, 0.1$ ;  $n = 5$ ) (Fig. 1d; individual values in Table S1). Conversely, PTC GL-3 inclusions increased (mean [SD],  $0.3$  [ $0.94$ ]; 95% CI:  $-0.6, 1.2$ ;  $n = 7$ ) during that period in those receiving placebo. Upon switching from placebo to migalastat at month 6, PTC GL-3 inclusions decreased by a mean (SD) of  $-0.7$  ( $0.91$ ) (95% CI:  $-1.7, 0.2$ ;  $n = 6$ ) over the next 6 months in these patients. These results mirror those reported for the overall amenable FACETS population, in which PTC GL-3 inclusions decreased over 6 months of migalastat treatment (mean [SE],  $-0.25$  [ $0.10$ ];  $n = 25$ ) and increased (mean [SE],  $0.07$  [ $0.13$ ];  $n = 20$ ) over 6 months of placebo treatment.<sup>20</sup> In the overall amenable FACETS population, a subsequent mean (SE) decrease in mean GL-3 inclusions per PTC of  $-0.33$  ( $0.15$ ;  $n = 17$ ) was seen following switch from placebo to migalastat at month 6.

Decreases in mean (SD) number of PTC GL-3 inclusions were seen in other patients whether treated with migalastat from month 0 to 6 ( $-0.1$  [ $0.30$ ]; 95% CI:  $-0.2, 0.0$ ;  $n = 20$ ) or switched from placebo to migalastat at month 6 (change from month 6 to 12;  $-0.1$  [ $0.24$ ]; 95% CI:  $-0.3, 0.1$ ;  $n = 11$ ) (Fig. 1d). A small decrease was also noted in other patients treated with placebo from month 0 to 6 ( $-0.05$  [ $0.10$ ]; 95% CI:  $-0.1, 0.0$ ;  $n = 13$ ). Overall, the decreases in the other patients subgroup were small, likely due to low baseline levels; however, the magnitude of change with migalastat was twice that of placebo.

#### GL-3 in glomerular cells

On the basis of qualitative assessments of kidney biopsy samples, patients in the classic phenotype subgroup had either a reduction or no change in GL-3 inclusions in glomerular endothelial cells (3/9 decreased, 6/9 had no change), mesangial cells (3/9 decreased, 6/9 had no change), and podocytes (9/9 had no change) at month 12 compared with baseline or month 6 (for patients switched from placebo to migalastat). No patient showed increase in GL-3 in these cell types after 6–12 months of migalastat treatment (Table S2).

A substantial portion of patients in the other patients subgroup had a reduction in GL-3 in glomerular cells, including mesangial cells (14/31), endothelial cells (8/31), and podocytes (6/31) at month 12. One patient had an increase in GL-3 in mesangial cells. The remaining patients experienced no change in GL-3 inclusions in these cells after 6–12 months of migalastat treatment.

#### Plasma lyso-Gb<sub>3</sub>

Migalastat was associated with a reduction in plasma lyso-Gb<sub>3</sub> in both subgroups, with a considerable reduction observed in

the classic phenotype subgroup. The average (SD) change from baseline to month 24 in plasma lyso-Gb<sub>3</sub> was  $-36.8$  ( $35.78$ ) nmol/L (95% CI:  $-69.9, -3.7$ ;  $n = 7$ ) for males with the classic phenotype and  $-7.66$  ( $20.27$ ) nmol/L (95% CI:  $-16.6, 1.3$ ;  $n = 22$ ) for other patients (Fig. 1e; individual values in Table S1). Notably, mean (SD) baseline lyso-Gb<sub>3</sub> levels were  $99.8$  ( $35.28$ ;  $n = 7$ ) nmol/L in the classic phenotype subgroup versus  $29.3$  ( $48.32$ ;  $n = 24$ ) nmol/L in the other patients subgroup.

#### PBMC $\alpha$ -Gal A activity

Within the classic phenotype subgroup, 6 months of randomized treatment with migalastat led to a mean (SD) increase in PBMC  $\alpha$ -Gal A activity of  $2.2$  ( $3.0$ ) 4 MU/h/mg (95% CI:  $-0.5, 5.0$ ;  $n = 7$ ), whereas treatment with placebo led to a substantially smaller increase ( $0.1$  [ $0.1$ ] 4 MU/h/mg; 95% CI:  $0.0, 0.2$ ;  $n = 7$ ). Increase in PBMC  $\alpha$ -Gal A activity was sustained during open-label migalastat; mean (SD) change from baseline to month 24 was  $2.6$  ( $2.3$ ) 4 MU/h/mg (95% CI:  $0.8, 4.4$ ;  $n = 9$ ). Changes in PBMC  $\alpha$ -Gal A activity appeared to be more variable in the other patients subgroup due to the small number of male patients; mean (SD) change from baseline to month 24 was  $2.7$  ( $9.8$ ) 4 MU/h/mg (95% CI:  $-21.7, 27.1$ ;  $n = 3$ ).

#### Adverse events

The most frequently reported adverse event that occurred during migalastat treatment was headache in both the classic phenotype (4/14 [29%]) and other patients (15/36 [42%]) subgroups. The other patients subgroup did not report any other adverse events that occurred in  $\geq 20\%$  of patients, whereas diarrhea, procedure pain, and vertigo were reported in  $\geq 20\%$  of patients in the classic phenotype subgroup (Table S3).

## DISCUSSION

Fabry disease is associated with a wide spectrum of disease progression and clinical phenotypes, ranging from the most severe, classic phenotype to an asymptomatic phenotype in a subset of female patients.<sup>1</sup> Newborn screening studies estimate the frequency of the classic phenotype as up to 1 in 22,570 males.<sup>27</sup> Patients with the classic phenotype typically have onset of symptoms during childhood or adolescence, followed by multiorgan complications that can progress to organ failure.<sup>1</sup> Therefore, early treatment initiation is crucial in this patient population. Current treatment guidelines and consensus recommendations recommend ERT treatment for all males  $\geq 16$  years old with classic Fabry disease, even if they are asymptomatic, presenting a high treatment burden.<sup>28</sup>

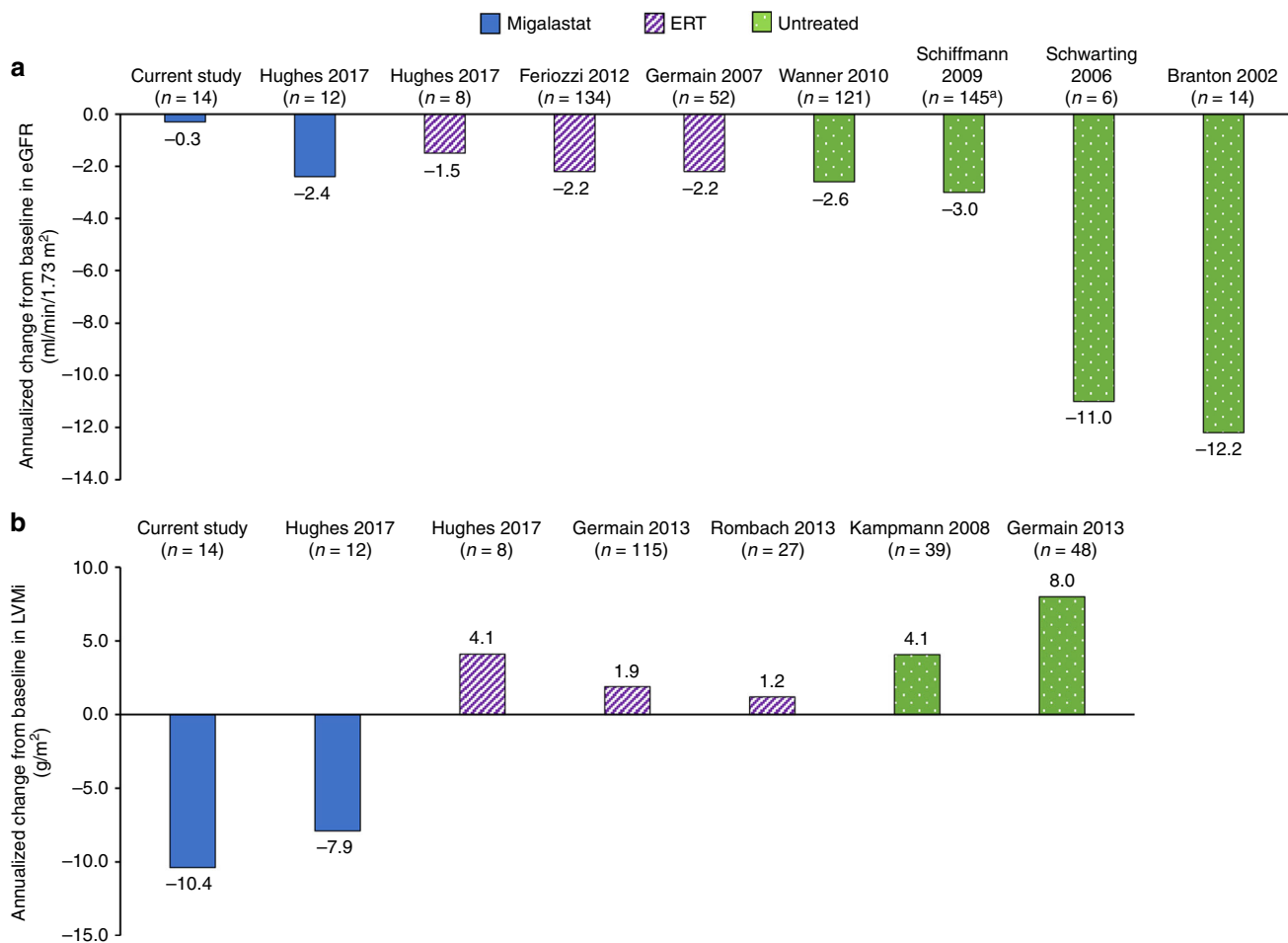
In a previously published report from the phase 3 FACETS study, patients with Fabry disease who have amenable variants were shown to benefit from treatment with migalastat.<sup>20</sup> The results presented here provide additional evidence for the beneficial effects of migalastat in male patients with the classic phenotype—the most severe form of Fabry disease.



In vitro assay data suggest that all patients with Fabry disease and amenable variants, including those with the classic phenotype, would benefit from migalastat treatment, as migalastat increases  $\alpha$ -Gal A activity in the lysates of cells expressing mutated enzyme.<sup>4</sup> This hypothesis is supported by our data, which demonstrate reduction in cardiac mass, improvements in GSRs-D, stabilization of  $eGFR_{CKD-EPI}$ , reduction in PTC GL-3 inclusions, plasma lyso-Gb<sub>3</sub>, and PBMC  $\alpha$ -Gal A activity in male patients with the classic phenotype treated with migalastat for up to 24 months.

Compared with published data for male patients with Fabry disease, the results of this analysis suggest that patients treated with migalastat had better renal<sup>29–32</sup> and cardiac<sup>33,34</sup> outcomes than that historically reported for untreated patients. Historical studies of untreated males report an annualized change in  $eGFR$  of up to  $-12.2$  mL/min/1.73 m<sup>2</sup>.<sup>29–32</sup> Comparatively, the annualized change in  $eGFR_{CKD-EPI}$  was  $-0.3$  mL/min/1.73 m<sup>2</sup> in the subgroup of patients with the classic phenotype treated with migalastat in FACETS (Fig. 2a). For LVMi, studies of untreated males report annualized

increases ranging from 4.1 to 8.0 g/m<sup>2</sup> (refs.<sup>33,34</sup>), compared with an annualized change of  $-10.4$  g/m<sup>2</sup> with migalastat in males with the classic phenotype (Fig. 2b). Furthermore, migalastat appears to provide potential cardiac benefit compared with ERT (Fig. 2) (refs.<sup>12,19,34–36</sup>). This is further supported by the greater efficacy of migalastat versus ERT in reducing cardiac mass at month 18 in a phase 3, open-label, randomized clinical trial (ATTRACT; NCT01218659) (ref.<sup>19</sup>). The benefit of migalastat on cardiac structure may reflect the broad tissue distribution of migalastat as an orally administered small molecule, including penetration into cardiac tissue.<sup>19,23</sup> However, the mechanism by which migalastat reduces LVMi is unknown. Clearance of GL-3 deposits from cardiomyocytes and vascular endothelial cells may have contributed to the mass reduction; GL-3 clearance from the latter could also lead to enhanced regional myocardial perfusion and function, influencing the dynamic of left ventricular remodeling. Additionally, it appeared that reduction in LVMi was due to a reduction in both wall thickness and cavity size (Amicus Therapeutic data on file).



**Fig. 2 Comparisons of published data for male patients with Fabry disease.** Annualized changes in (a) renal function and (b) LVMi as reported in the literature for male patients who are either untreated or have been treated with migalastat or ERT. <sup>a</sup>Change from baseline calculated using the weighted average for end-stage renal disease subgroups ( $n = 128$  males with non-end-stage renal disease and  $n = 17$  males with end-stage renal disease).  $eGFR$  estimated glomerular filtration rate,  $ERT$  enzyme replacement therapy,  $LVMi$  left ventricular mass index.

Further investigations are needed to obtain more definitive understanding of the pathophysiology of cardiomyopathy in Fabry disease and the effects of migalastat on cardiac tissues.

Elevated plasma lyso-Gb<sub>3</sub> levels have become recognized as a diagnostic indicator for Fabry disease, especially for classic Fabry patients.<sup>37</sup> Consistent with previous reports,<sup>37,38</sup> baseline plasma lyso-Gb<sub>3</sub> levels were highly elevated in male patients with the classic phenotype in FACETS. Migalastat was associated with substantial reduction in lyso-Gb<sub>3</sub> in this patient subgroup. Despite the reduction, plasma lyso-Gb<sub>3</sub> levels remained elevated above the normal ranges after 24 months of treatment.<sup>38</sup> It has been reported that lyso-Gb<sub>3</sub> levels decreased but did not normalize despite years of ERT in male patients with the classic phenotype.<sup>37</sup>

In our study, the classic phenotype was defined as males with multiorgan system involvement and residual PBMC  $\alpha$ -Gal A activity <3% of normal (Table 1). Of note, one of the *GLA* variants associated with the classic phenotype, p.Arg301Gln (p.R301Q), was originally identified in a Japanese male and characterized as a cardiac variant, a milder atypical variant with late-onset cardiac manifestations limited to the heart.<sup>39</sup> Since then, at least one case of a more severe phenotype has been associated with this variant, including a 45-year-old male patient who had end-stage renal disease in addition to cardiomyopathy; this patient benefited from double heart and kidney transplant.<sup>40</sup> The patient with the p.Arg301Gln variant in our study had multiorgan disease (stage 3 chronic kidney disease [eGFR 56 mL/min/1.73 m<sup>2</sup>], abnormal LVMI [137 g/m<sup>2</sup>]) and PBMC  $\alpha$ -Gal A activity 1.9% of WT, thus fulfilling the criteria for inclusion in the classic phenotype subgroup.

Our study includes a descriptive analysis of the baseline disease characteristics and clinical outcomes of two patient subgroups in FACETS; the other patients subgroup consists predominantly, though not exclusively, of females. One potential limitation of our study is that outcomes based on indirect comparisons across studies should be interpreted with caution because of potentially confounding factors, such as variations in study design, inclusion/exclusion criteria, baseline demographics, and disease severity. In addition, the results reported for males with the classic phenotype from the FACETS study are derived from post hoc analyses in a relatively small group of patients observed for up to 24 months, a relatively short study duration for long-term data, given the progressive nature of Fabry disease. Lastly, GRS-D has not been validated specifically in Fabry disease, although it has demonstrated reliability, stability, and construct validity across a range of GI disorders. No GI symptom rating scale has been validated in Fabry disease. Despite these limitations, our data show beneficial effects in male patients with the classic phenotype, and provide support for the use of migalastat in all patients with amenable *GLA* variants regardless of disease severity.

In conclusion, migalastat, a first-in-class oral pharmacologic chaperone, has previously demonstrated clinical benefits in patients with Fabry disease and amenable *GLA* variants.<sup>20</sup> The

data reported here establish that migalastat benefited male patients with the classic phenotype. During the phase 3 FACETS trial, migalastat treatment increased endogenous  $\alpha$ -Gal A activity, stabilized eGFR, reduced LVMI, improved diarrhea symptoms, and reduced PTC GL-3 inclusions and plasma lyso-Gb<sub>3</sub> in male patients with migalastat-amenable *GLA* variants leading to the classic phenotype of Fabry disease.

## SUPPLEMENTARY INFORMATION

The online version of this article (<https://doi.org/10.1038/s41436-019-0451-z>) contains supplementary material, which is available to authorized users.

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## DISCLOSURE

D.P.G. has received honoraria and research grants from Amicus, Sanofi Genzyme, and Shire. K.N. has served as an advisor for Amicus, Shire, and Sanofi Genzyme, has received research support from Amicus and Shire, and has received travel support from Sanofi Genzyme. R.G. has received honoraria from Amicus, BioMarin, Sanofi Genzyme, and Shire. D.G.B. has received research funding, serves as a consultant, and is on the speaker's bureau for Amicus and Sanofi Genzyme, and has received research funding from Shire. D.A.H. has served as a consultant for and received research funding and honoraria from Amicus, Shire, Sanofi Genzyme, Protalix, and Actelion. L.M.B. has served as a consultant for Protalix. R.B.C. has served as a consultant for Amicus and has received grants from the National Institutes of Health. N.S., J.P.C., E.B., and J.A.B. are employees of and hold stock in Amicus. At the time the work was done, C.V. was an employee and held stock in Amicus. J.C.J. declares no conflicts of interest.

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