

Proof of Mechanism Studies with Bitopertin, a Selective Glycine Transporter 1 Inhibitor Under Development for the Treatment of Erythropoietic Protoporphyrria (EPP) and X-linked Protoporphyrria (XLPP)

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INTRODUCTION

Bitopertin is a selective and orally available inhibitor of glycine transporter 1 (GlyT1), which supplies extracellular glycine for the initial step of heme biosynthesis in erythroid cells. Bitopertin has been studied extensively in clinical trials (>4,000 human subjects), has a favorable safety profile, and has been shown to modulate the heme biosynthesis pathway. Disc Medicine is developing bitopertin as a first-in-class, potentially disease modifying therapy for EPP and XLPP, two rare genetic disorders of heme biosynthesis. EPP is caused by a partial deficiency of the enzyme ferrochelatase (FECH), the last enzyme in the heme biosynthesis pathway that incorporates iron into protoporphyrin IX (PPIX). In EPP patients, a loss of >65% of FECH activity results in accumulation of PPIX, which leads to painful cutaneous photosensitivity with erythema and edema and hepatobiliary injury. In XLPP, mutations in the C-terminal domain of 5'-Aminolevulinatase Synthase 2 (ALAS2), the first enzyme in heme production, result in a gain of function leading to elevated 5-aminolevulinic acid (5-ALA) production, and PPIX accumulation (Fig. 1A). We hypothesize that, by inhibiting glycine uptake into erythroid precursors, bitopertin can restrict heme pathway metabolite production and reduce the disease-causative PPIX accumulation in the blood of EPP and XLPP patients (Fig. 1B).

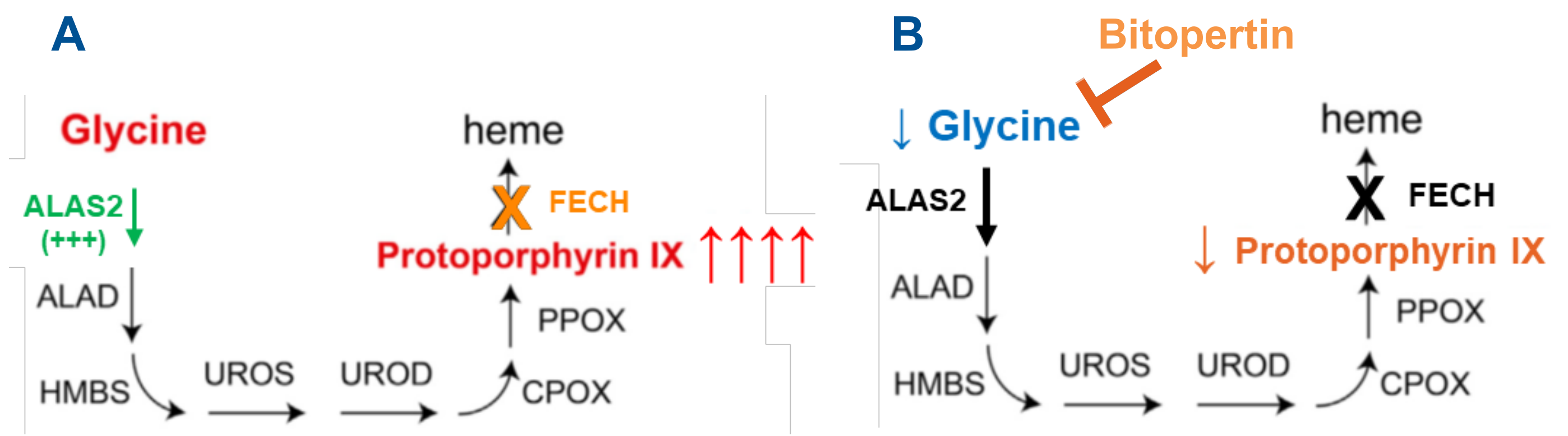


Figure 1A. Enzyme defects in EPP and XLPP lead to the accumulation of PPIX, a phototoxic metabolite

Figure 1B. Bitopertin, a GlyT1 inhibitor, reduces PPIX accumulation in EPP and XLPP

OBJECTIVES

- To establish cellular models of EPP using human erythroleukemia K562 and human cord blood CD34+ cells
- To evaluate the effect of bitopertin on reducing PPIX accumulation in EPP cellular models
- To evaluate the effect of bitopertin on PPIX accumulation and hemoglobin levels in mouse models of EPP and XLPP

METHODS

- For K562 cells, we performed CRISPR-Cas9 genome editing to knock down one FECH allele and introduce the FECH c.315-48C hypomorphic variant of the EPP genotype in trans.
- For CD34+ cells, knock down (>60% reduction) of FECH mRNA was achieved with lentiviral vectors expressing shRNA of FECH. Transduced cells were differentiated for 9 days with Bitopertin or ORG-25543.
- The efficacy of bitopertin to treat EPP was further evaluated in female *Fech^{m1Pas}* EPP mouse model and in male *Alas2^{Q548X/Y}* XLPP mouse model. The recessive *Fech^{m1Pas}* allele is an ethylnitrosourea (ENU)-induced missense mutation that retains approximately 5% residual ferrochelatase activity (Tutois et al, J Clin Invest. 1991; 88: p1730). The *Alas2^{Q548X}* animals were generated by employing CRISPR-Cas9 editing technology to introduce a known human gain-of-function mutation (S. Ducamp et al, poster). *Fech^{m1Pas}* and *Alas2^{Q548X}* mice were fed with diet containing 0 or 100 ppm bitopertin for 8 weeks starting at 6 weeks of age. Effects on PPIX and hemoglobin were determined at the end of 8 weeks of dosing.

RESULTS

Establishment of K562 cellular model of EPP

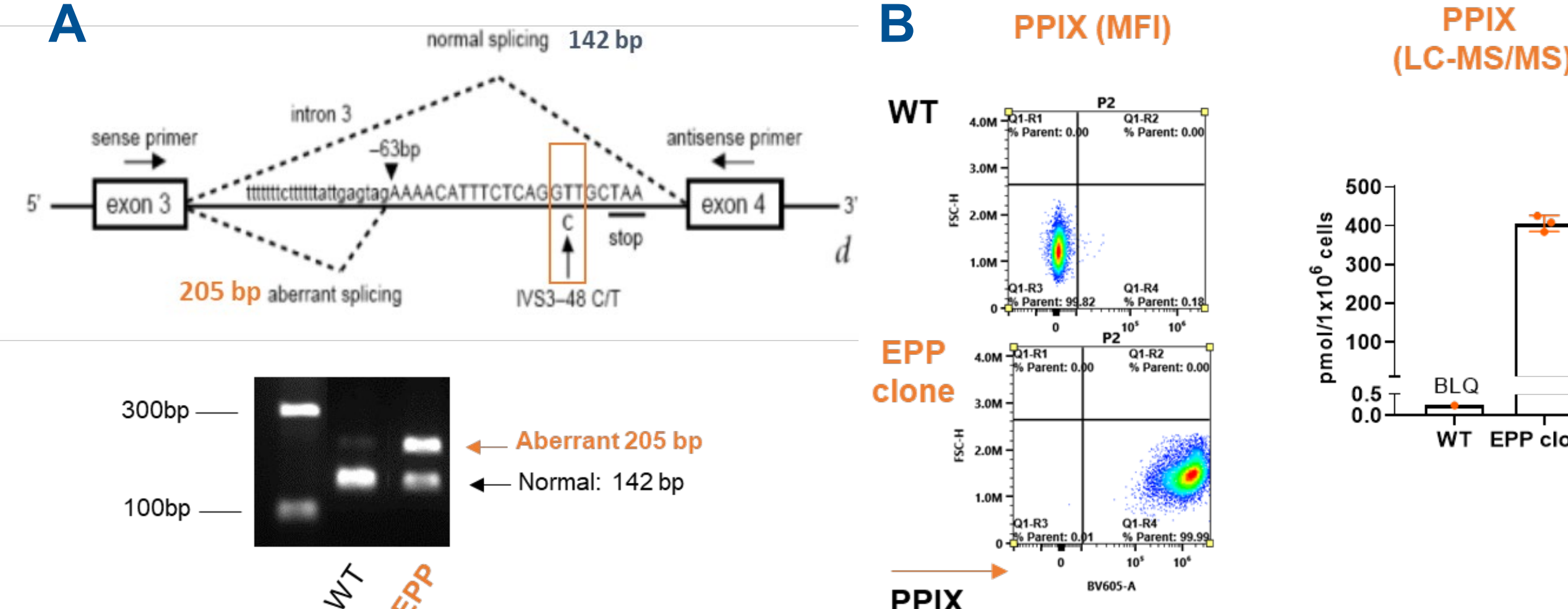


Figure 2A. The intronic IVS3-48T>C mutation increases the use of an aberrant splice site

Figure 2B. IVS3-48C/KO mutations resulted in supra-physiological levels of PPIX

Compound	K562 EPP assay (EC ₅₀ , nM)	GlyT1 uptake assay (IC ₅₀ , nM)
Bitopertin	9	25
PF-03463275	46	12
ALX-5407	0.34	3
ORG-24598	5.6	7
LY-2365109	4.1	16
ORG-25543 (GlyT2 inhibitor)	no inhibition	>10,000

* Literature values

Table 1. Activity of known GlyT1 and GlyT2 specific inhibitors in the K562 cellular model of EPP

Bitopertin reduced PPIX in K562-EPP cell model

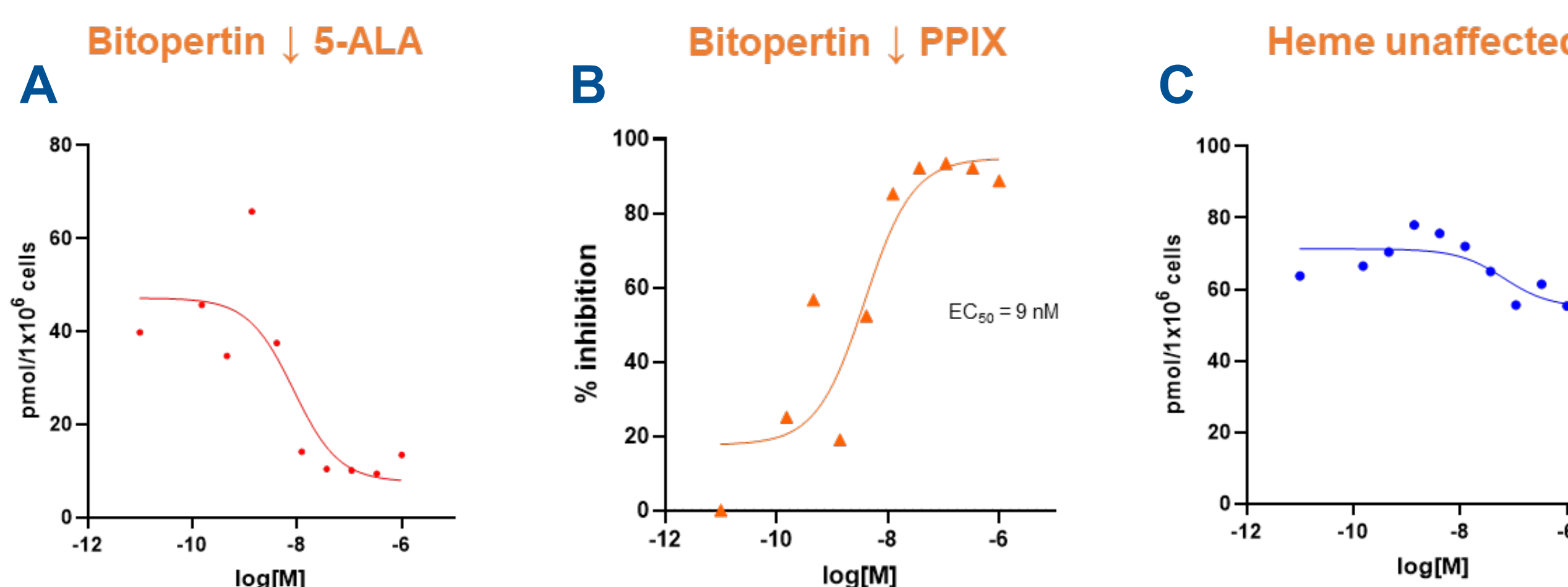


Figure 3. Bitopertin decreased formation of 5-ALA (A), prevented PPIX accumulation in a dose dependent manner (B), without affecting heme formation (C)

Bitopertin reduced PPIX in hCD34+ model of EPP

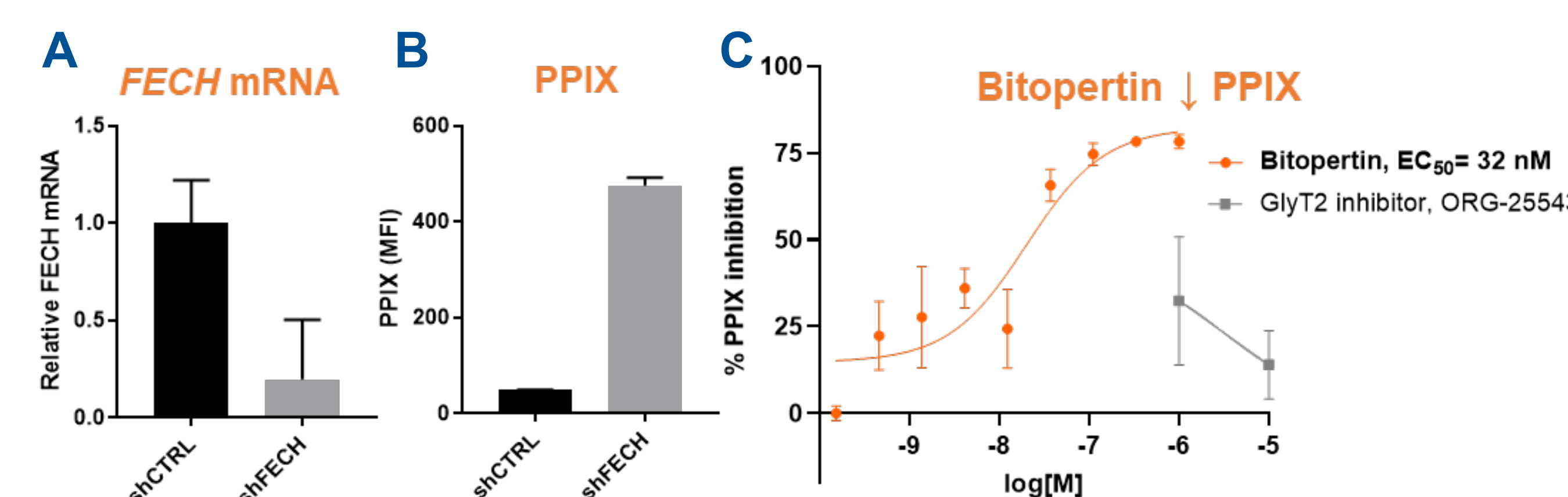


Figure 4. ~75% reduction in FECH mRNA level observed (4A); leading to PPIX accumulation in cells (B)

Figure 4C. Bitopertin reduced PPIX with an EC₅₀ of 32 nM, while GlyT2 inhibitor ORG-25543 had minimal effect

Bitopertin reduced PPIX in *Fech^{m1Pas}* mice

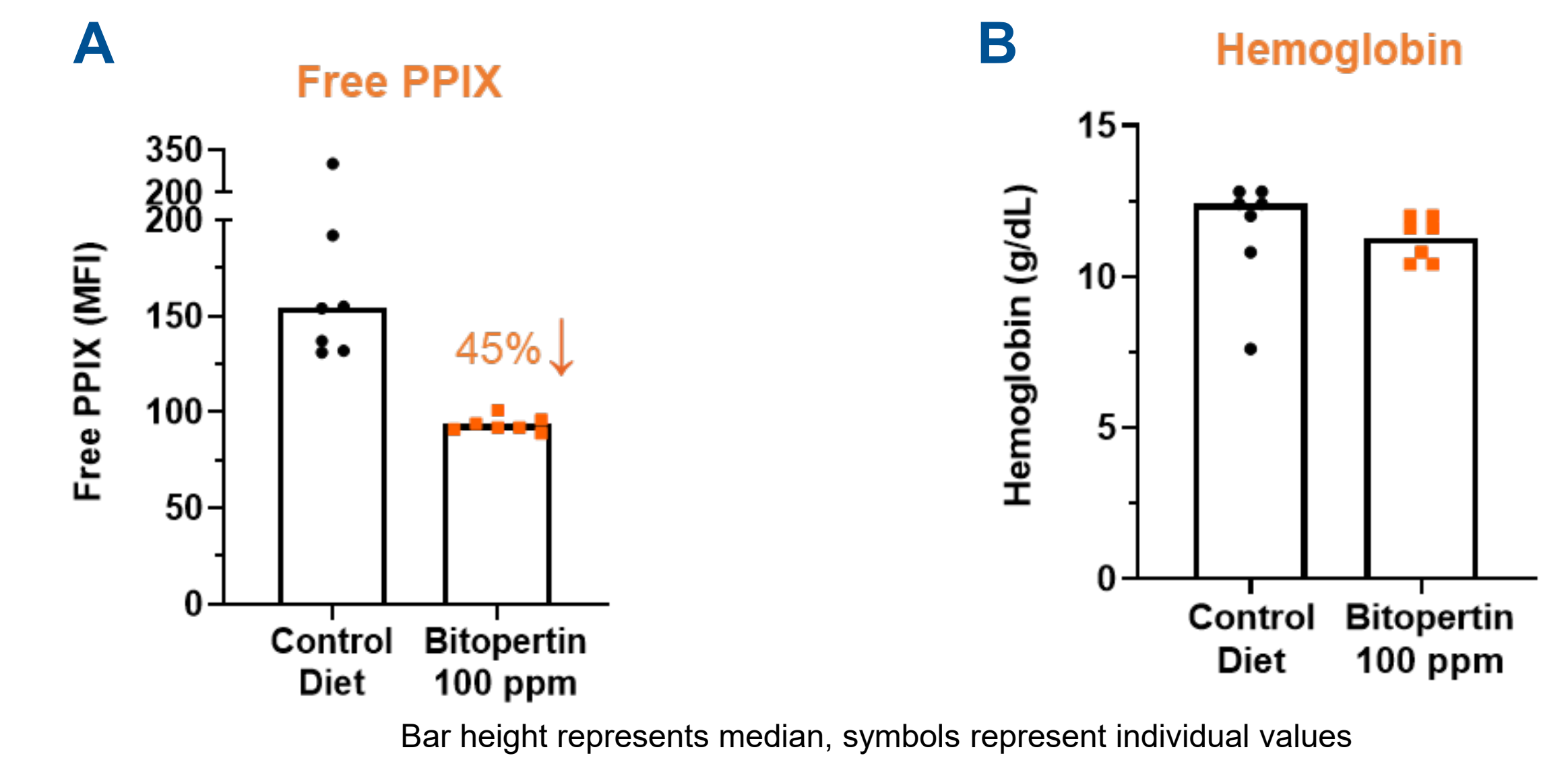


Figure 5. Effects of bitopertin in EPP mouse model (*Fech^{m1Pas}* / *Fech^{m1Pas}*) on PPIX (A) and hemoglobin levels (B) after 8 weeks of treatment on 100ppm bitopertin diet

Bitopertin reduced PPIX in *Alas2^{Q548X}* mice

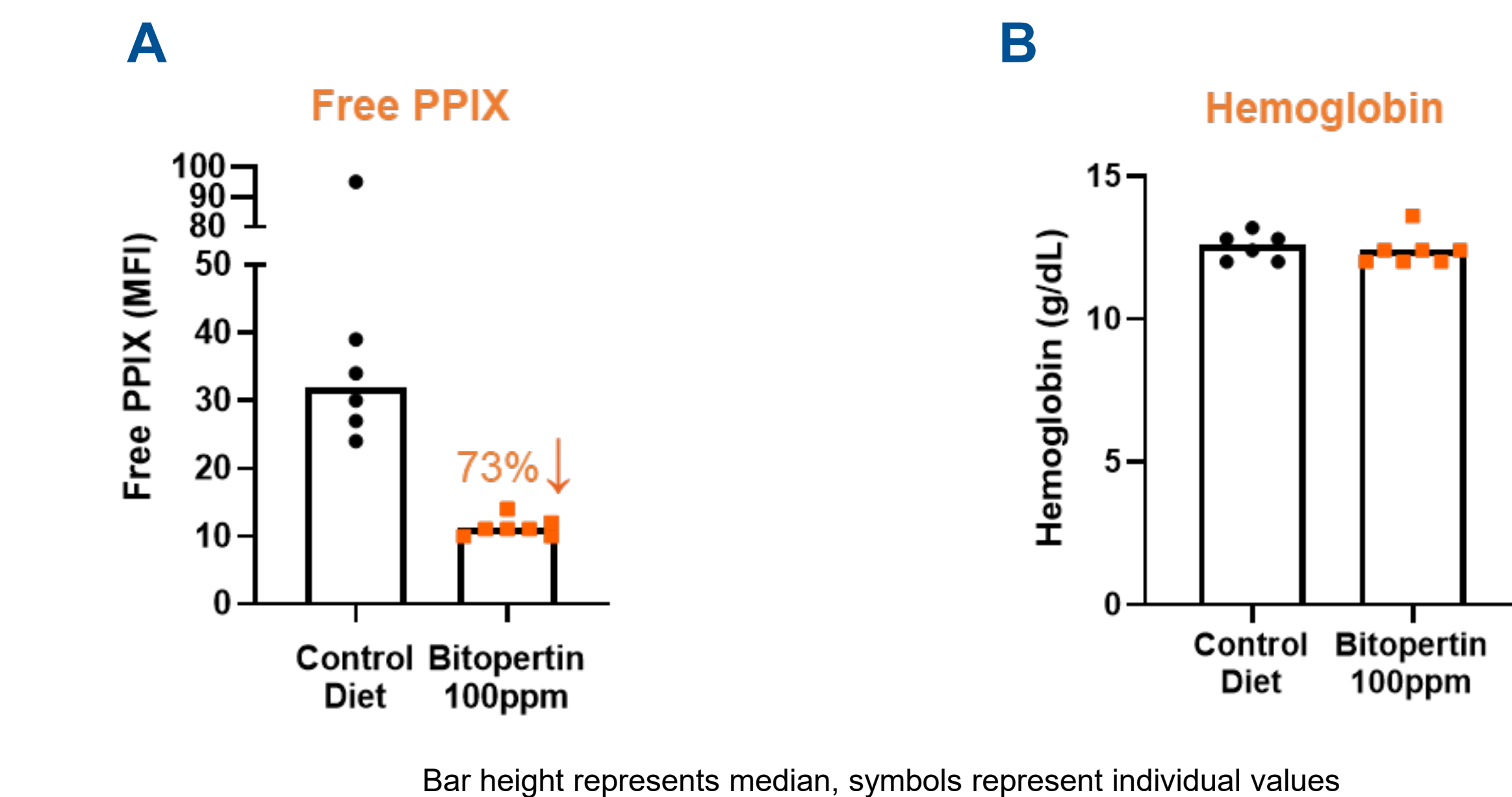


Figure 6. Effects of bitopertin in XLPP mouse model (*Alas2^{Q548X/Y}*) on PPIX (A) and hemoglobin levels (B) after 8 weeks of treatment on 100ppm bitopertin diet

CONCLUSIONS

- Bitopertin is a selective GlyT1 inhibitor with a well-characterized safety profile in humans
- Bitopertin reduced PPIX in K562 and hCD34+ cellular models of EPP
- Bitopertin reduced PPIX in mouse models of EPP and XLPP without effects on hemoglobin. Target reduction of 30-50% exceeded.
- Bitopertin has the potential to improve light tolerance and hepatobiliary injury in EPP patients by decreasing PPIX in erythrocytes

CONTACT INFORMATION

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