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## A phase 2 study of HQK-1001, an oral fetal haemoglobin inducer, in $\beta$ -thalassaemia intermedia

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#### Keywords

Anaemia; β-thalassaemia; clinical trials; gene expression; fetal haemoglobin; haemoglobinopathies

The  $\beta$ -thalassaemia syndromes represent a World Health Organization-designated global health burden (Weatherall *et al*, 2010). Reactivation of fetal globin (HbF) expression is a rational therapeutic approach in inherited  $\beta$ -globindisorders, because the fetal *HBG*( $\gamma$ -globin genes) are universally present and appropriately contextually integrated in the *HBB*( $\beta$ -globin gene) locus in haematopoietic stem cells (Bauer *et al*, 2012). The defective production of  $\beta$ globin chains in  $\beta$ -thalassaemia can be compensated for by an increase in  $\gamma$ -globin chains, which pair with  $\alpha$ -globinchains to form HbF, thereby decreasing the  $\alpha$ /non  $\alpha$ -globin imbalance, the hallmark of  $\beta$ -thalassaemia. Several classes of HbF inducers have been investigated in  $\beta$ -thalassaemia, including cytotoxic agents, DNA methyl transferase inhibitors, histone deacetylase (HDAC) inhibitors including short chain fatty acids, thalidomide derivative, and erythropoietin, but no consistently effective agents have been identified (Musallam *et al*, 2013).

The short chain fatty acids arginine butyrate, sodium phenylbutyrate, and isobutyramide were shown to increase HbF in  $\beta$ -thalassaemia and sickle cell disease, but had to be administered intravenously or by large oral daily doses, which is not practical for widespread long-term use (Perrine *et al*, 1993; Collins *et al*, 1995; Capellini *et al*, 2000). The orally bio available butyrate derivative 2,2-dimethylbutyrate sodium salt (HQK-1001) does not exhibit HDAC2 inhibitory activity and stimulates *HBG* expression and erythropoiesis in animal models and *in vitro* at concentrations readily achievable in humans(Pace *et al*, 2002; Mankindy *et al*, 2006). In a proof-of-concept study, HQK-1001 at

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Authorship: AI, SP, and RG designed the research; AI, MK, AT, SK and TA performed research; DC and HA performed genetic studies; RG and SP analysed the data and wrote the manuscript; AI and SP revised the manuscript.

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10, 20, 30, and 40 mg/kg administered daily for eight weeks in 21 subjects with nontransfusion dependent  $\beta$ -thalassaemia was well-tolerated (Fuchareon *et al*, 2013). HQK-1001 at 20 mg/kg, which provided the best results, increased HbFin 8 of 9 subjects with a median increase of 6.6% and 4.4 g/l, and increased total haemoglobin in 4 of 9 subjects by a mean of 11 g/L. Here, a single-centre study was conducted to evaluate HQK-1001 at 20 mg/kg/day administered for a longer period (NCT01642758).

Adult patients with  $\beta$ -thalassaemia intermedia characterized by two  $\beta$ -globin mutations were eligible if their haemoglobin was between 60 and 90 g/l on two occasions during the 30-day screening period. Patients were excluded if they were transfused within the previous three months, received iron chelation agents within the previous seven days, another investigational agent within the previous 30 days, erythropoietic agents within the previous 90 days, or hydroxycarbamide within the previous six months, or had pulmonary hypertension requiring oxygen therapy, alanine aminotransferase (ALT) > 4 times the upper limit of normal, or serum creatinine > 135 µmol/l. HQK-1001 capsules (HemaQuest Pharmaceuticals, San Diego, CA) was administered at 20 mg/kg once daily for 24 weeks. Folic acid was given daily, and to prevent iron-deficient inefficient erythropoiesis, oral iron was given if serum ferritin was < 1500 pmol/l, but stopped if ferritin levels were > 2250 pmol/l. After signing an Ethics Committee approved informed consent form, subjects were assessed clinically and underwent laboratory tests twice during a 30-day screening period, every four weeks while receiving HQK-1001, and then four weeks after the end of dosing.

Ten subjects were enrolled, seven male and three female, with a mean age of 29.4 years (range 18-52 years). Eight subjects were splenectomized; two had palpable splenomegaly at 4 and 7 cm below the left costal margin. The mean (range) baseline values were: HbF26.6% (7.9–73.8%), absolute HbF 20.1 g/l (5.5–53.9 g/l), total haemoglobin 77.4 g/l (61.5–96.0 g/l), platelet count  $782 \times 10^9/l$  (486-1039  $\times 10^9/l$ ), reticulocytes 10.9% (7.1–15.7%), and serum ferritin 3188 pmol/l (375–9772 pmol/l).

Nine subjects completed the study and one subject was discontinued at Week 16 because of worsening anaemia requiring a transfusion. Mean compliance with HQK-1001, calculated as the ratio of the number of HQK-1001 capsules taken divided by the number of capsules prescribed, was 92.5%; two subjects had compliance <90%. Treatment was generally well-tolerated. All adverse events, except one case of vertigo, were graded as mild or moderate, and were reversible. Fatigue was the most common adverse event, reported in 3 subjects. In contrast, 5 subjects reported increased activity and improved mood. Two subjects each reported nausea, epigastric pain, dyspepsia or fever. The most common laboratory abnormalities were mild and reversible increases in aspartate aminotransferase (AST) in five subjects and in ALT in four.

HbF increased in all subjects, with peak increase occurring after a mean of 14 weeks of therapy; the mean (range) increase from baseline was 4.8% (2.3–9.8%) for HbF % (p = 0.0006) and 3.19 g/l (0.5–6.6 g/l) for absolute HbF (p = 0.001). Total haemoglobin increased in 7 subjects, with a mean increase of 4.7 g/l (range 1.0–10.0 g/l). Figure 1 shows the baseline and peak value by subject for HbF and total haemoglobin. Table I presents each subject's thalassaemia mutations and polymorphisms for 3 quantitative trait loci (QTL) that were shown to strongly influence baseline HbF levels (Thein *et al*, 2009). Seven subjects were homozygous for the IVS I-6 (C-T)  $\beta$ + thalassemia mutation and only 3 were heterozygous for a favourable genetic modifier.

This study demonstrates that HQK-1001 at 20 mg/kg/day for 24 weeks was well tolerated, significantly increased HbF, and modestly increased total haemoglobin. An interim analysis of a recently completed study of HQK-1001 at 20 mg/kg/day for 26 weeks in 10 patients

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with Hb E- $\beta$ -thalassaemia showed higher mean increase in HbF of 10% (range 4.3–20.9%), with an increase in total haemoglobin > 5 g/l in 3 subjects (Fuchareon *et al*, 2012). These patients all had a  $\beta^{O}$ -thalassaemia mutation, and 9 had at least one favorable allele for the Xmn-I QTL, which is linked to the *HBB*:c.79G>A( $\beta^{E}$  globin) gene in that population. Three trials have now demonstrated that HQK-1001 increases HbF in  $\beta$ -thalassaemia. It remains to be determined whether the magnitude of increase in HbF is sufficient to reduce long-term complications of chronic haemolysis, ineffective erythropoiesis, anaemia, and transfusion requirements. Further studies of genetically characterized patients for longer periods appear warranted.

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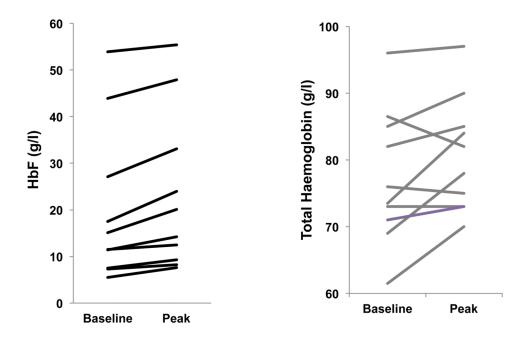


Figure 1. Baseline and peak values for HbF and total haemoglobin

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Table I

# **Baseline Characteristics**

Thalassaemia Mutations	HMIP <sup>*</sup> Rs9399137	BCLIIA F
IVS I-6/IVS I-6 (T>C)	(N/N)	(N/

Subject ID		Age (Years) Baseline HbF %	Thalassaemia Mutations	HMIP* Rs9399137	BCL11A Rs766432	HBG2Xmn-1 Rs7482144
001	23	20	IVS I-6/IVS I-6 (T>C)	(N/N)	(N/+)	(N/N)
002	39	7.9	IVS I-6/IVS I-6 (T>-C)	(N/N)	(N/N)	(N/N)
003	32	20	IVS I-110 G>A, IVS II-1G>A	(N/N)	(+/+)	(N/+)
004	21	58	IVS I-6/IVS I-110	(N/N)	(N/+)	(N/N)
005	36	25	IVS I-6/IVS I-6 (T>C)	(H/+)	(N/N)	(N/N)
008	25	20	9-I SAI/9-I SAI	(N/N)	(N/N)	(N/N)
600	52	17	(T>C)CD29/CD29	(N/+)	(N/N)	(N/N)
011	27	9.8	IVS I-6/IVS I-6 (T>C)	(N/N)	(N/N)	(N/N)
012	25	15	IVS I-6/IVS I-6 (T>-C)	(H/+)	(N/N)	(N/N)
013	18	54	IVS I-6/IVS I-6 (T>-C)	(H/+)	(N/N)	(N/N)
For rs9399137,	, N designates the	e major allele, T; + de	For rs9399137, N designates the major allele, T; + designates the minor allele, C.			

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For rs 766432, N designates the major allele, A; + designates the minor allele, C.

\* HMIP = *HBS1L*-*MYB* intergenic polymorphism.