# SYNB1353, A Proposed Therapy for Homocystinuria, Lowers Plasma Methionine and Homocysteine in Healthy Volunteers Exposed to a Methionine Challenge

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## Introduction

Classical homocystinuria (HCU) is an inherited disorder caused by pathogenic variants in the cystathionine beta-synthase (CBS) gene **(Figure 1)** resulting in excessive accumulation of homocysteine (Hcy) and multiorgan clinical manifestations. Early initiation of methionine-restricted diets significantly lowers the risk of developing complications in HCU. Elevated Hcy levels are associated with impairments of the eye, skeletal system, vascular system, and central nervous system.<sup>1</sup> In patients with residual CBS activity (~50% of HCU population), vitamin B6 (pyridoxine) is effective at reducing Hcy levels. For pyridoxine unresponsive patients, betaine (involved in remethylation of Hcy to methionine) and a low-methionine diet<sup>2</sup> that is very low in natural protein are the current therapeutic options.

## **Study Design**

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We evaluated SYNB1353 in a double-blinded, placebo-controlled Phase I trial utilizing a multiple ascending dose (MAD) design (Figure 3A). Four cohorts using dose levels,  $3x10^{11}$ ,  $6x10^{11}$  and  $1x10^{12}$  SYNB1353 live cells, were evaluated for safety, tolerability, and capacity to metabolize methionine in four cohorts of healthy volunteers challenged with 30 mg/kg methionine before and after exposure to SYNB1353. Study demographics are shown in Table 1. Subjects received a methionine challenge on day -1, followed by SYNB1353 using a dose ramp on days 1 to 7 (Figure 3B). The methionine challenge was repeated on day 7 before SYNB1353 dosing. Changes in plasma methionine and total Hcy

## **Results (continued)**

Methionine Load Leads to Increase in Plasma Methionine and Plasma Homocysteine in Healthy Volunteers

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Figure 1. Mutations in the CBS gene cause the accumulation of Hcy and HCU. In HCU

were assessed over 24h after the methionine challenge.



**Figure 3. Phase 1 study design and dosing schedule. (A)** Each cohort (n=8) included six subjects dosed with SYNB1353 and two subjects dosed with placebo. Safety and tolerability were assessed after each cohort prior to dose escalation. The 1x10<sup>12</sup> live cell cohort was repeated with an alternate formulation. (B) In each cohort, SYNB1353 or placebo was administered over a seven-day period with meals starting with a single dose on day 1, two doses on day 2 and 3, and three doses on days 4 to 7. A methionine challenge (30 mg/kg) was administered on day -1 and day 7 and plasma methionine and plasma total Hcy were measured over 24h.

#### Time post-methionine load (hours)

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**Figure 4. Increased plasma methionine and total Hcy following methionine challenge. (A)** Plasma methionine and **(B)** total Hcy levels of all subjects (n=30) from all cohorts over 24 h after receiving a 30 mg/kg methionine load challenge. Data represent the mean ± SD. Methionine challenge resulted in an 11-fold increase in plasma methionine ( $C_{max}$ ) and a 2-fold increase in plasma total Hcy ( $C_{max}$ ) compared to baseline values.

## Proof-of-Mechanism: SYNB1353 Blocks Methionine Absorption and Lowers Plasma Methionine



patients, mutations in the *CBS* gene result in accumulation of Hcy. Pharmacotherapeutic options for the treatment of HCU consist of vitamin B6 (pyridoxine), which can lower Hcy levels in B6-responsive patients, and betaine, which is involved in Hcy remethylation to methionine

#### SYNB1353: A methionine metabolizing synthetic biotic

SYNB1353 was engineered from the probiotic *E. coli* Nissle (EcN) to metabolize methionine (Met) via the methionine decarboxylase (MetDC) pathway, preventing its conversion into homocysteine. SYNB1353 converts Met to 3-methylthiopropylamine (3-MTP). To prevent the release of methionine once it enters the cell, the YjeH gene was deleted which is responsible for Met export in EcN.



#### Table 1. Study demographics

	Placebo	3×10 <sup>11</sup>	6×10 <sup>11</sup>	1×10 <sup>12</sup>	1×10 <sup>12*</sup>	
Parameter	(N=8)	(N=6)	(N=5)	(N=6)	(N=5)	
Age (years)						
Mean (SD)	37.9 (8.4)	39.7 (12.8)	32.8 (4.7)	33.2 (11.3)	47.4 (14.0)	
Min, Max	26, 47	25, 56	25, 36	23, 54	31,63	
Sex [F/M]	0/8	3/3	1/4	2/4	2/3	
Race						
African American	4	4	2	0	5	
White	4	2	3	6	0	
Other	0	0	1	0	0	
Ethnicity						
Hispanic	1	1	1	2	2	
Not Hispanic	7	5	4	4	3	
BMI (kg/m²)						
Mean (SD)	26.9 (3.7)	29.0 (4.2)	26.8 (4.6)	25.9 (3.4)	27.5 (3.4)	
Min, Max	22.16, 32.70	24.56, 34.80	21.82, 33.65	21.07, 30.72	23.95, 31.83	
Completed Dosing n (%)	8 (100)	5 (83.3)	5 (100)	6 (100)	5 (100)	
Discontinued n (%)	0	1 (16.7)	0	0	0	

## Results

### SYNB1353 was generally well-tolerated in healthy volunteers

Dose	Dose	Dose	Dose	Total	rate per	rate p

**Figure 5. SYNB1353 blocks methionine absorption.** Percent change from baseline in day 7 (A) plasma methionine  $AUC_{0-24h}$  and (B) total plasma Hcy  $AUC_{0-24h}$  for cohorts receiving  $1\times10^{12}$  live cell of SYNB1353. FORM = formulation. LS mean change, 95% CI \*p< 0.05

#### **SYNB1353 Metabolizes Methionine**



Cohort	SYNB1353 Dose	% Positive Samples	Mean Ae (nmol)
Placebo		50% (4/8)	1139
	3x10 <sup>11</sup>	17% (1/6)	1330
SYNB1353	6x10 <sup>11</sup>	20% (1/5)	5303
	1x10 <sup>12*</sup>	73% (8/11)	4865

#### Predose Placebo SYNB1353

**Figure 6. 3MTP-glycine is increased in SYNB1353 dosed subjects.** Total amount of excreted (Ae) 3-MTP-glycine in urine was determined for all subjects pre-dose on day -1 (n=30), all placebo dosed subjects (n=8) and all subjects dosed with SYNB1353 (n=22). Urine was collected for 24h after methionine challenge on either day -1 (predose) or on day 7. Data represent the mean and SD of all positive samples collected (n=4/30 predose, n=4/8 placebo and n=10/22 SYNB1353). The table shows the number and percentage of positive samples in each cohort and the corresponding mean. \*pooled data from both cohorts receiving  $1 \times 10^{12}$  live cells.

## Conclusions

SYNB1353 was **well tolerated in healthy volunteers** with GI adverse event rates and severity similar between active and placebo groups

**Figure 2.** Schematic of SYNB1353. Met enters through the importer MetP into the cell and is converted via MetDC to 3-MTP. The strain contains a deletion of *yjeH*, a methionine exporter. 3-MTP is further metabolized *in* vivo to 3-MTP-glycine (3-MTP-gly) by the liver thus preventing further conversion to Hcy. Measurement of 3-MTP-gly provides evidence of strain activity *in vivo* 

#### **References:**

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	3E+11	6E+11	1E+12	1E+12	SYNB1353	subject	Placebo	subject
Cohort	(n=6)	(n=5)*	(n=6)	(n=5)*	(n=22)	SYNB1353	(n=8)	Placebo
Subjects with at least one TEAE	3	1	4	3	11	0.50	3	0.38
Maximum Grade 1	0	1	0	3	4	0.18	0	0.00
Maximum Grade 2	3	0	4	0	7	0.32	3	0.38
Total Number of TEAE	16	2	17	3	38	1.73	9	1.13
Subjects with at least one IMP-related TEAE	3	0	5	3	11	0.50	1	0.13
Maximum Grade 1	1	0	0	1	2	0.09	1	0.13
Maximum Grade 2	2	0	5	2	9	0.41	0	0.00
Any SAE	0	0	0	0	0	0.00	0	0.00
TEAE causing withdrawal	0	0	0	0	0	0.00	0	0.00

**Table 2. Adverse events in healthy subjects receiving SYNB1353 or placebo.** The majority of TEAE were GI related in nature. Rates and severity of TEAE were similar between SYNB1353 and placebo groups.

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SYNB1353 has demonstrated methionine metabolism in the GI tract of
healthy volunteers, resulting in a lowering of plasma methionine and
production of 3MTP-glycine, assessed following a meal challenge to elevate
methionine levels
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Previously presented mechanistic modeling data suggests that SYNB1353 may increase protein intake and **lower plasma Hcy by up to 58% in HCU patients**<sup>3</sup>

Based on this proof of mechanism in healthy volunteers, **SYNB1353 will be advanced to a Phase 2** proof of concept study in patients with HCU

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