

A Novel Treatment Approach For Treating OTC Deficiency By Targeting RegRNAs Using Oligonucleotides

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CAMP4 is pioneering the field of regulatory RNAs to upregulate genes



regRNAs are key controllers of gene expression



In the nucleus, genes and their regulatory elements are organized into conserved 3D DNA structures known as <u>Insulated</u> <u>Neighborhoods</u> to <u>control gene expression</u>.

regRNAs are uniquely transcribed within neighborhoods and act as <u>rheostats for</u> precise genomic control.

<u>CAMP4</u> has created the only <u>platform to</u> <u>drug regRNAs.</u>

RNA Actuating Platform charts a proprietary path from RNA targets to drugs



Next-gen sequencing powered by proprietary AI to identify regRNAs



Screen RNA Actuators

Screen oligo drug candidates to target regRNA hotspots for maximum gene upregulation



Program for Druggability

Design high-potency RNA Actuators for safe and effective delivery to target tissues underlying disease





regRNA sequence AAUUCACUU<mark>UUAAUACAGCUCUG</mark>GAGUGGUGG Chemistry variations



Elevating residual OTC activity by at least 2-fold addresses 90% of patients

OTC Deficiency: largest subsegment of UCD patient population (50%)

- Primary function of Urea cycle is detoxification of ammonia to urea for elimination
- Disease severity and age of onset varies with residual OTC activity and ammonia levels; strong genotype-phenotype relationship



are critical for ammonia → urea conversion

Therapeutic goal: increase OTC gene product to reach >5% of normal

Individual OTC-deficient Patient Data¹



OTC gene is located on X chromosome

1: Wakiya et al., Mol Gen Met, 2012 (105)-404

CAMP4 platform identified novel regRNA transcripts to upregulate OTC mRNA



Overall process of RNA Actuator screening

Total 406 sequences were screened for targeting OTC regRNAs





regRNA targeting RNA Actuators upregulate OTC mRNA dose-dependently



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RNA actuators led to increased ureagenesis in patient cells¹

OTC mRNA levels in response to treatment

Ureagenesis read-out cultured cells





P values calculated by one-way ANOVA

- c.-106C>A (Allele ID 480410, late-onset OTC deficiency) pathogenic (dbSNP: rs749748052), leading to decreased OTC mRNA¹.
- Variant associated with 10-25% of normal OTC activity

Developing a mouse surrogate for Otc mRNA

Identified regRNA for mouse Otc

The human *OTC* enhancer is structurally • conserved in mouse but the sequences are not conserved

Designed and screened RNA actuators

Screened for impact on Otc mRNA with • **RNA** Actuators

Otc deficient (Otc^{spf/ash}) mouse model¹

- The spf ^{ash} mouse has a variant c.386G>A, p.Arg129His in the Otc gene that impact splicing
- Otc mRNA levels are decreased (5~12%) of wt control) in spf/ash livers
- Male spf ^{ash} mice have a mild biochemical phenotype with low OTC activity (5%-10%) of wild-type)

RNA Actuators increase mouse *Otc* mRNA levels



To summarize...

- Identified regRNA that controls OTC gene expression
- Identified regRNA targeting RNA actuators that can upregulate human OTC mRNA in a dose-dependent manner -> increase in ureagenesis
- RNA actuators targeting mouse regRNA increase Otc mRNA in both wildtype and Otc deficient cells
- This approach offers a novel way of treating a disease caused by a hypomorphic allele by upregulating the endogenous gene expression using RNA Actuators.
- Our results indicate that CAMP4's RAP[™] technology may provide precise, potent therapeutics that can be
 programmed to treat thousands of diseases.

<u>Upcoming</u>

- Test lead human OTC regRNA targeting RNA Actuators for
 - With GalNAc to test for NHP pharmacology, safety study
- Test lead mouse Otc regRNA targeting RNA Actuators for
 - In Otc deficient mice for Proof of Concept (PoC) read-out