DAYBREAK Trial: Setmelanotide vs placebo in patients with melanocortin-4 receptor pathway variants

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ECO 2025 - Conflict Of Interest

Name: Erica van den Akker

□ I have the following potential conflicts of interest to report:

X Research Contracts

X Consulting

Employment in the Industry

□ Stockholder of a healthcare company

□ Owner of a healthcare company

Other(s)

□ I declare that I have no potential conflict of interest.

The central hypothalamus is a key regulator of energy balance, appetite, and bodyweight

• The hypothalamic melanocortin-4 receptor (MC4R) pathway is a key regulator of energy balance and food intake^{1,2}



AgRP, agouti-related peptide; LEPR, leptin receptor; MC4R, melanocortin-4 receptor; MSH, melanocyte-stimulating hormone; PCSK1, proprotein convertase subtilisin/kexin type 1; POMC, proopiomelanocortin. **1.** Farooqi IS. *Biol Psychiatry*. 2022;91:856–59; **2.** Yeo GSH, et al. *Mol. Metab*. 2021;48:101206.

Impaired MC4R signalling due to rare genetic variants may result in hyperphagia and obesity from the first years of life^{1–4}

- The MC4R agonist setmelanotide is approved by the FDA and EMA for the treatment of certain MC4R pathway diseases, controlling hyperphagia^{*} and treating obesity^{5,6}
- DAYBREAK (NCT04963231) was a two-stage Phase 2 study designed to evaluate the efficacy of setmelanotide in individuals carrying confirmed variants in ≥1 of 31 genes with strong or very strong relevance to the MC4R pathway⁷



*Absence of satiety and pathologic, insatiable hunger accompanied by abnormal food-seeking behaviors.

AgRP, agouti-related peptide; EMA, European Medicines Agency; FDA, Food and Drug Administration; LEPR, leptin receptor; MC4R, melanocortin-4 receptor; MSH, melanocyte-stimulating hormone; PCSK1, proprotein convertase subtilisin/kexin type 1; POMC, proopiomelanocortin.

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Objectives of this study

- To determine the efficacy of setmelanotide in achieving weight loss in children and adults with obesity related to gene variants selected by Rhythm's ClinGen-based framework based on their strong association with the MC4R pathway¹
- To identify which populations may have the best potential to benefit from setmelanotide therapy



1. Vogel et al. Presented at: European Society for Paediatric Endocrinology; September 22–26, 2021; Virtual.

Inclusion criteria and primary endpoint

Stage 1 eligibility criteria

- Genetic confirmation in patients
 6–65 years
- Patients with obesity
 - Adults (≥18 years): BMI ≥40 kg/m²
 - Paediatric (<18 years): BMI ≥97th percentile for age and sex

Primary endpoint

 Proportion of patients by genotype who achieved a BMI reduction of ≥5% from baseline at the end of Stage 1



Stage 2 eligibility criteria

- Reduction at end of Stage 1, from baseline
 - Adult: reduction of \geq 3% BMI
 - Paediatric: reduction of ≥3% BMI
 <u>OR</u> ≥0.05 BMI z-score

BMI, body mass index.

Study design

• Efficacy outcome assessed at end of Stage 2 was change from baseline in BMI (kg/m²)



*Virtual visit. BMI, body mass index; R, randomisation.

Patient disposition



Patient and demographics

• After Stage 1, 49 patients with PHIP, PLXNA(1-4), SEMA3(A-G), SIM1, MAGEL2, TBX3, or RPGRIP1L variants were randomised 2:1 to receive setmelanotide or placebo

Stage 2 cohort demographics			
	All	<18 y	≥18 y
Male, n (%)	22 (44.9)	10 (45.5)	12 (54.5)
Female, n (%)	27 (55.1)	14 (51.9)	13 (48.1)
	Mean (SD)	Range	% of Stage 1 starters (n/N)
BMI, kg/m ²			
Adult baseline (n=25)	46.1 (7.2)	40.4–69.9	23 (25/109)
Adult Stage 2 start (n=25)	42.6 (7.0)	36.2–66.3	-
BMI z-score (CDC)			
Paediatric baseline (n=24)	2.5 (0.3)	1.83–2.97	44 (24/55)
Paediatric Stage 2 start (n=24)	2.25 (0.4)	1.48-2.92	-

BMI, body mass index; CDC, Centers for Disease Control and Prevention; MAGEL2, melanoma-associated antigen gene L2; PHIP, pleckstrin homology domain interacting protein; PLXNA1-4, plexin A1-4; RPGRIP1L, retinitis pigmentosa GTPase regulator-interacting protein-1 like; SD, standard deviation; SEMA3A-G, semaphorin 3A-G; SIM1, single-minded homolog 1; TBX3, T-box transcription factor.

Results



A higher proportion of patients in the setmelanotide arm achieved or maintained 5% BMI reduction from study baseline to the end of Stage 2 Mean (SD; range) percent BMI change in the continuous setmelanotide arm from baseline to the end of Stage 2







- Results from patients with variants in MAGEL2 (n=1 adult; n=2 children) and RPGRIP1L (n=1 adult) were consistent: patients randomised to setmelanotide **continued or maintained weight loss** in Stage 2
- The patient with a TBX3 variant (n=1 child) was assigned to placebo, regained weight, received rescue setmelanotide open-label therapy, and **resumed losing weight**

BMI, body mass index; MAGEL2, melanoma-associated antigen gene L2; RPGRIP1L, retinitis pigmentosa GTPase regulator-interacting protein-1 like; SD, standard deviation; TBX3, T-box transcription factor.

Results in patients with PHIP variants

 Individuals with PHIP variants, which can lead to obesity by disrupting POMC transcription, maintained a consistent weight loss response



BMI, body mass index; PHIP, pleckstrin homology domain interacting protein; POMC, proopiomelanocortin.

Results in patients with SEMA3E variants

- SEMA3E had excellent response, but was limited by the sample size (n=3)
- SEMA3 (A D, F, G) had a more variable degree of weight loss



*One adult and 1 paediatric patient with a *SEMA3G* variant dropped out of Stage 2 before having any data and are not shown. BMI, body mass index; SEMA3A–G, semaphorin 3A–G.

Results in patients with PLXNA and SIM1 variants

• PLXNA (1-4) and SIM1 had a more variable degree of weight loss, some of substantially greater magnitude than PHIP



BMI, body mass index; PHIP, pleckstrin homology domain interacting protein; PLXNA1-4, plexin A1-4; SIM1, single-minded homolog 1.

Safety

• Setmelanotide was well tolerated with no new safety concerns across the entire study



Conclusion

- Clinical response to setmelanotide treatment, a highly selective MC4R agonist, suggests that the MC4R pathway may be a **key biologic driver** of obesity in patients with variants of interest
- The design of the exploratory DAYBREAK trial enabled the **identification of multiple genes of interest** that merit further investigation, by utilising
 - Stage 1 open-label run-in period to identify patients with impaired MC4R signalling
 - Stage 2 confirmation of response in the randomised withdrawal period
- The **percent change in BMI** from baseline to the end of Stage 2 varied between gene cohorts
- Further studies may elucidate whether the genetic variants of interest explored in DAYBREAK contribute to a loss of function in the MC4R pathway or can identify other patient-specific factors that can modulate response to setmelanotide treatment

BMI, body mass index; MC4R, melanocortin-4 receptor.