

# CMR outcomes following one-time *in vivo* CRISPR editing with nexiguran ziclumeran in patients with ATTR-CM: phase 1 results

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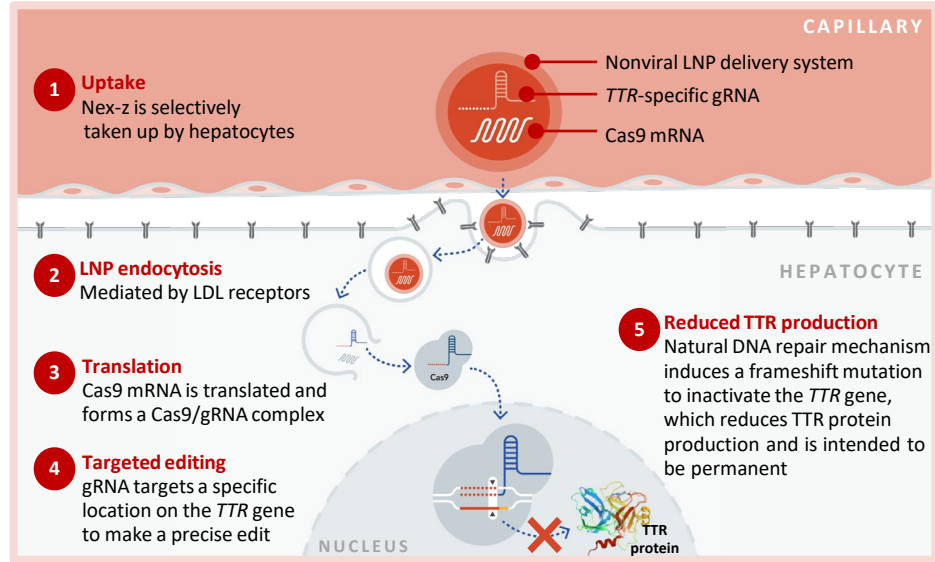
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11 May 2026

# Background and purpose

- ATTR-CM is characterised by progressive cardiac remodelling and dysfunction, decline in functional capacity and QoL, frequent hospitalization, and death<sup>1</sup>
- Cardiovascular magnetic resonance (CMR), including extracellular volume (ECV) mapping, provides a sensitive method to quantify amyloid burden and treatment response<sup>2,3</sup>
  - Higher ECV is a strong and independent predictor of mortality, with risk rising 22% for every 10% absolute increase in ECV<sup>3</sup>
- In a Phase 1 trial (NCT04601051) of patients with ATTR-CM, a single dose of nexiguran ziclumeran (nex-z) was associated with rapid, deep, and durable TTR knockdown, favourable biomarker and functional trajectories, and was generally well tolerated<sup>4</sup>

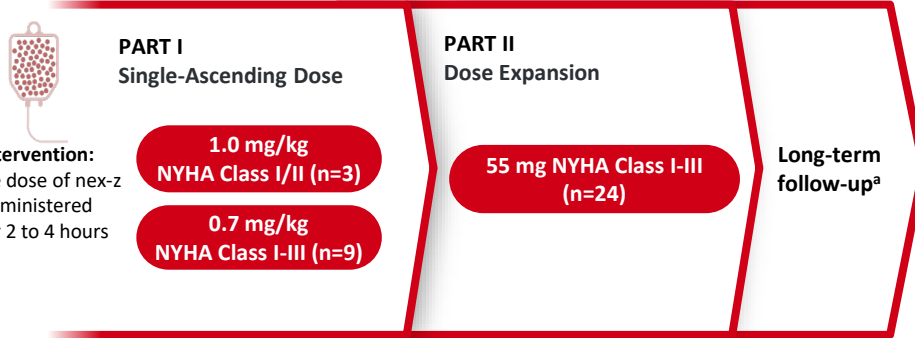
**Nex-z is an *in vivo* investigational one-time CRISPR/Cas9 therapy designed to inactivate the *TTR* gene, whether wild-type or variant<sup>1,5</sup>**



**Purpose:** Evaluate the impact of nex-z on cardiac structure, function, and myocardial amyloid load in patients with ATTR-CM

# Phase 1, two-part, open-label study of nex-z in adults with ATTR-CM

## Two-Part, Open-Label Study in Adults With ATTR-CM



### PRIMARY OBJECTIVES

Evaluate safety, tolerability, and PD

- Measure serum TTR levels

### SECONDARY OBJECTIVES

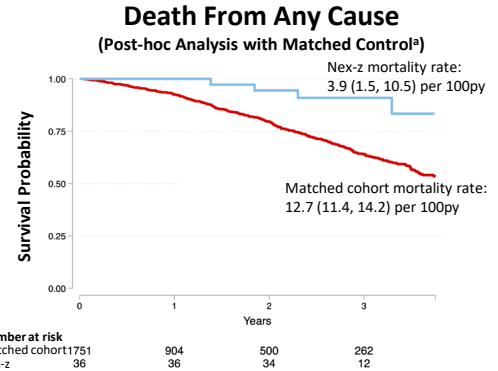
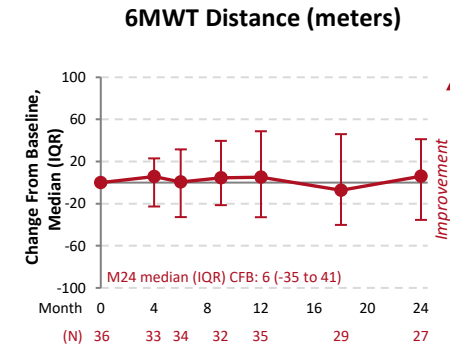
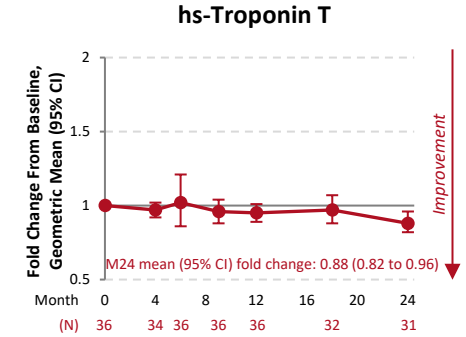
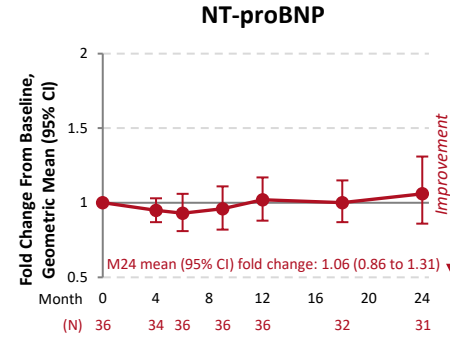
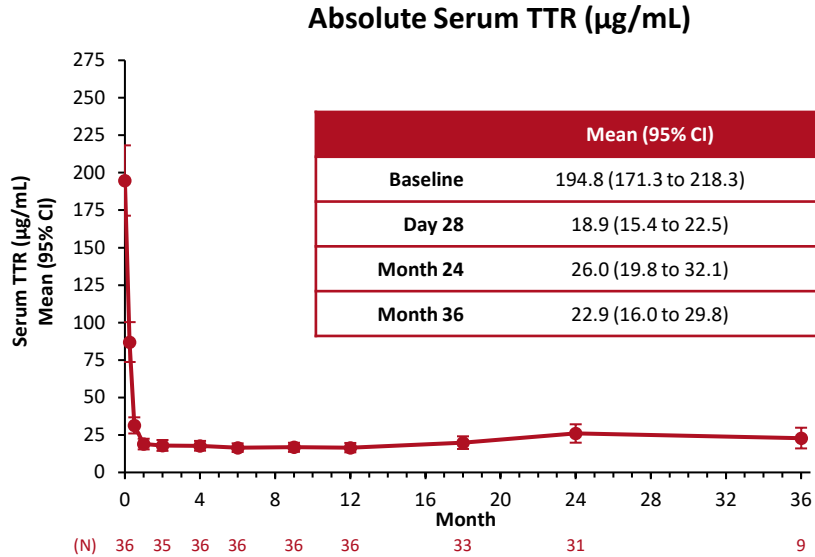
Evaluate efficacy on clinical measures of cardiac disease

- Biomarkers of disease progression, including NT-proBNP, hs-Troponin T, and 6MWT, cardiac imaging, and KCCQ score

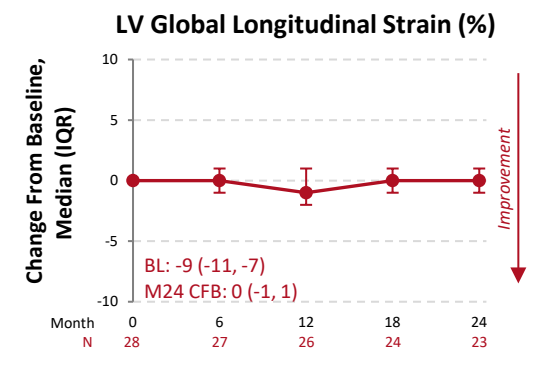
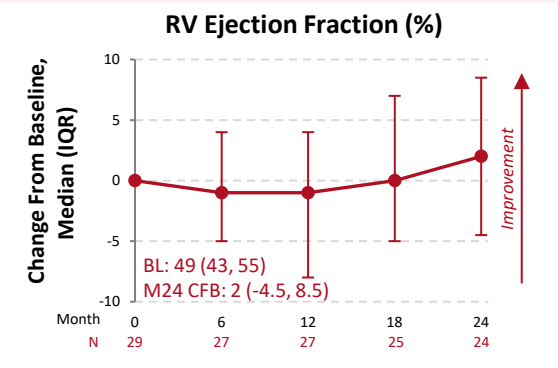
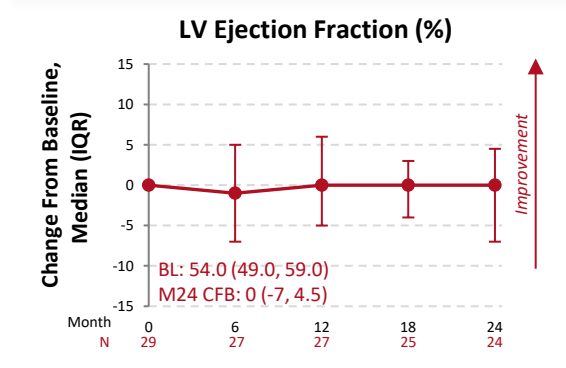
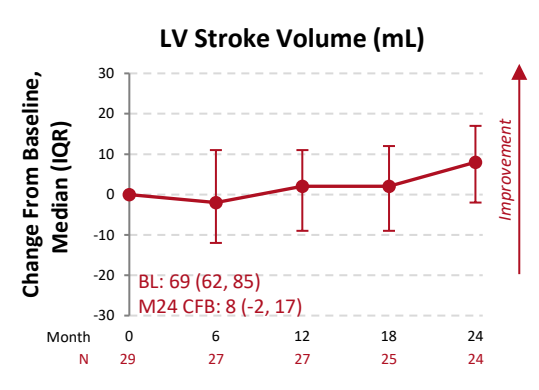
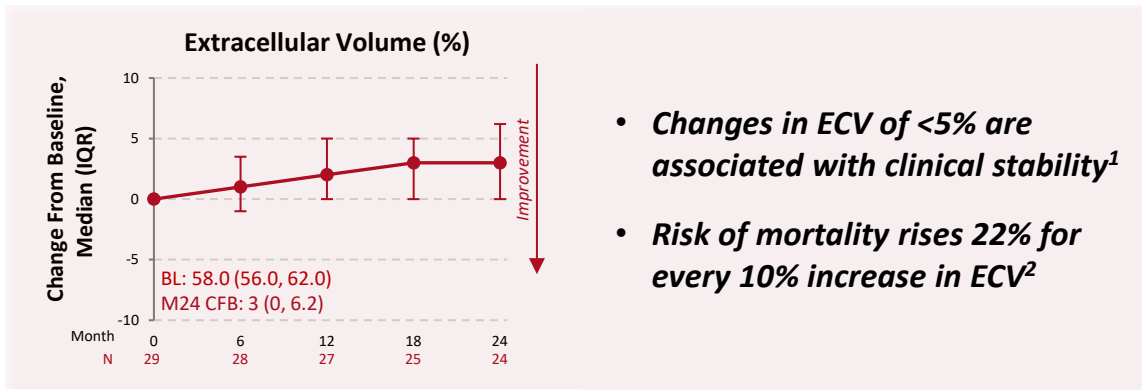
Characteristic	All patients (N=36)	
Age, median (min, max), y	78 (46, 90)	
Sex, male, n (%)	35 (97)	
Race, n (%)	Black or African descent	8 (22)
	White or Caucasian	28 (78)
NT-proBNP, median (min, max), ng/L	2052 (851, 19,624)	
hs-Troponin T, median (min, max), ng/L	56 (15, 204)	
eGFR, median (min, max), mL/min/1.73 m <sup>2</sup>	64 (30, 87)	
6MWT distance, median (min, max), m	331 (178, 580)	
Peak VO <sub>2</sub> , median (min, max), mL/kg/min	12.7 (7.8, 28.4)	
CMR extracellular volume, median (min, max), %	58 (45, 71)	
TTR genotype, n (%)	Wild type	25 (69)
	p.V142I <sup>b</sup>	7 (19)
	Other mutations	4 (11)
NYHA class, n (%)	I	3 (8)
	II	15 (42)
	III	18 (50)
Tafamidis use at baseline, n (%)	0 (0)	

## Population representative of ATTR-CM, including patients with advanced disease

# Nex-z led to durable reductions in serum TTR and stabilized or improved markers of cardiomyopathy and function in patients with ATTR-CM<sup>1</sup>



# Nex-z was associated with stability of cardiac structure, function, and amyloid burden over 24 months, as assessed by CMR<sup>a</sup>



# Safety and conclusions

Event	All patients (N=36) n (%)
<b>At least one AE</b>	36 (100)
<b>AEs occurring in ≥15% of patients</b>	
Cardiac failure	13 (36)
COVID-19	8 (22)
Upper respiratory tract infection	7 (19)
Atrial fibrillation	6 (17)
Urinary tract infection	6 (17)
<b>Treatment-related AEs in ≥5% of patients</b>	
Infusion-related reaction	5 (14)
Aspartate aminotransferase increased	2 (6)
<b>Any AE leading to treatment discontinuation</b>	0
<b>Any SAE<sup>a</sup></b>	15 (42)
<b>SAEs occurring in ≥5% of patients</b>	
Cardiac failure	8 (22)
Urinary tract infection	3 (8)
Acute myocardial infarction	2 (6)
Atrial flutter	2 (6)
Cardiac failure congestive	2 (6)
Pneumonia	2 (6)
<b>Any event leading to death<sup>b</sup></b>	4 (11)

## Conclusions

In this Phase 1 cohort with a high proportion of hereditary ATTR-CM, a single dose of nex-z was associated with stability of cardiac structure, function, and amyloid burden over 2 years.

These findings support the potential of gene editing to alter the natural history of amyloid deposition and disease progression in ATTR-CM.