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Reporting Summary

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

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For	all statistical ar	alyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.		
n/a	Confirmed			
	The exact	sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement		
\boxtimes	A stateme	ent on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly		
	The statis Only comm	tical test(s) used AND whether they are one- or two-sided non tests should be described solely by name; describe more complex techniques in the Methods section.		
	A descript	cion of all covariates tested		
	A descript	cion of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons		
	A full desc	cription of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) tion (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)		
\boxtimes		ypothesis testing, the test statistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted es as exact values whenever suitable.		
	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings			
	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes			
\boxtimes	\square Estimates of effect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated			
	'	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.		
So	ftware an	d code		
Poli	cy information	about <u>availability of computer code</u>		
Da	ata collection	Data was collected through the electronic data capture system Medidata Classic Rave 2019-2023		
Da	ata analysis	SAS Version 9.4 or higher and R version 3.5.2 or higher was used for analyses.		

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.

Data

Policy information about availability of data

All manuscripts must include a data availability statement. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

Data originating from the trial is considered commercially sensitive and as such, is not publicly available. To the extent that current legislation allows this, Tthe authors will provide access to individual-deidentified participant-level data that underlie the data presented in this paper, to researchers who provide a methodologically sound proposal for academic purposes to interpret, verify and extend research in the article that does not violate intellectual property or

confidentiality obligations beginning 12 months after article publication. Researchers should contact the corresponding author when applying for data access. Use of data will be restricted to the agreed upon purpose.

Research involving human participants, their data, or biological material

Policy information about studies with <u>human participants or human data</u>. See also policy information about <u>sex, gender (identity/presentation)</u>, and sexual orientation and race, ethnicity and racism.

Reporting on sex and gender

Subjects self-reported sex. No analyses by sex have been reported due to the limited sample size making such subgroup analyses difficult, if not impossible to interpret.

Reporting on race, ethnicity, or other socially relevant groupings

No data on race was collected in the trial and all subjects self-reported to be "Not Hispanic or Latino". No analyses by ethnicity was performed in the trial.

Population characteristics

Reported in Table 1

Recruitment

Subjects were recruited through hospitals in Hungary and Denmark. Subjects were required to have an recent onset episode of atrial fibrillation and were identified mostly in emergency rooms or through collaboration with general practitioners when they presented with an acute episode of atrial fibrillation. Subjects were approached by a health care professional and asked if they would be interested in participating in the trial.

Ethics oversight

The trial protocol was approved by the following ethics committees: Den Videnskabsetiske Komité, Region Sjælland, Alléen 15, 4180 Sorø, Denmark; Egészségügyi Tudományos Tanács Klinikai Farmakológiai Etikai Bizottsága, Széchenyi István tér 7-8, Budapest, Hungary.

Note that full information on the approval of the study protocol must also be provided in the manuscript.

Field-specific reporting

Please select the one be	elow that is the best fit for your research.	. If you	u are not sure, read the appropriate sections before making your selection.	
X Life sciences	Behavioural & social sciences		Ecological, evolutionary & environmental sciences	

For a reference copy of the document with all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

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The sample size was selected based on a similar previous trial (doi:10.1016/J.JACC.2004.09.021), and no formal sample size calculations were made.

Data exclusions

Sample size

Subjects were excluded from the primary analysis if they did not fulfill the following pre-specified definition of the full analysis set: randomised participants who were administered double-blind study treatment and have an evaluable AF conversion status within 90 min from the start of infusion.

The exclusion was done to ensure that only participants that could contribute data to the primary analysis were included in this. No data points were excluded in the analysis.

Replication

As this was a stand-alone clinical trial no replication was performed.

Randomization

Randomization was performed with the use of a computer-generated random-sequence and interactive voice- and web-response system

Blinding

The trial was double-blind, meaning that both subjects and investigators were blinded to treatment allocation.

Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

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Materials & experimental systems	Methods	
n/a Involved in the study	n/a Involved in the study	
Antibodies	ChiP-seq	
Eukaryotic cell lines	✓ ✓ Flow cytometry	
Palaeontology and archaeology	MRI-based neuroimaging	
Animals and other organisms	'	
Clinical data		
Dual use research of concern		
Plants		
'		
Clinical data		
Policy information about <u>clinical studies</u>		
All manuscripts should comply with the ICMJE <u>guide</u>	<u>lines for publication of clinical research</u> and a completed <u>CONSORT checklist</u> must be included with all submissions.	
Clinical trial registration The trial was pre-report NCT04571385	gistered in EudraCT: number 2018-004445-17. Later, it was also registered in ClinicalTrials.gov: number	
Study protocol Available in the Sup	orotocol Available in the Supplementary Appendix	
The trial was conducted at 10 hospital sites in Hungary and Denmark and patients were randomized between September 24, 2019, and December 9, 2022. Data was entered by site personal into an electronic data capture system (Medidate Rave Classic) and verified by sponsor monitors based on source documents.		
registries. The prima	The primary and secondary endpoints were prespecified in the trial protocol and in the Eudra CT and Clinicaltrials.gov public trial registries. The primary endpoint of cardioversion within 90 minutes was assessed by the investigator and verified centrally based on Holter monitoring read by a core laboratory. The secondary efficacy outcomes were assessed by the investigator and, if Holter	

this was used to verify the assessment.

monitoring was available at the assessment time (Holter monitoring was done for at least 8 and up to 24 hours after infusion start),