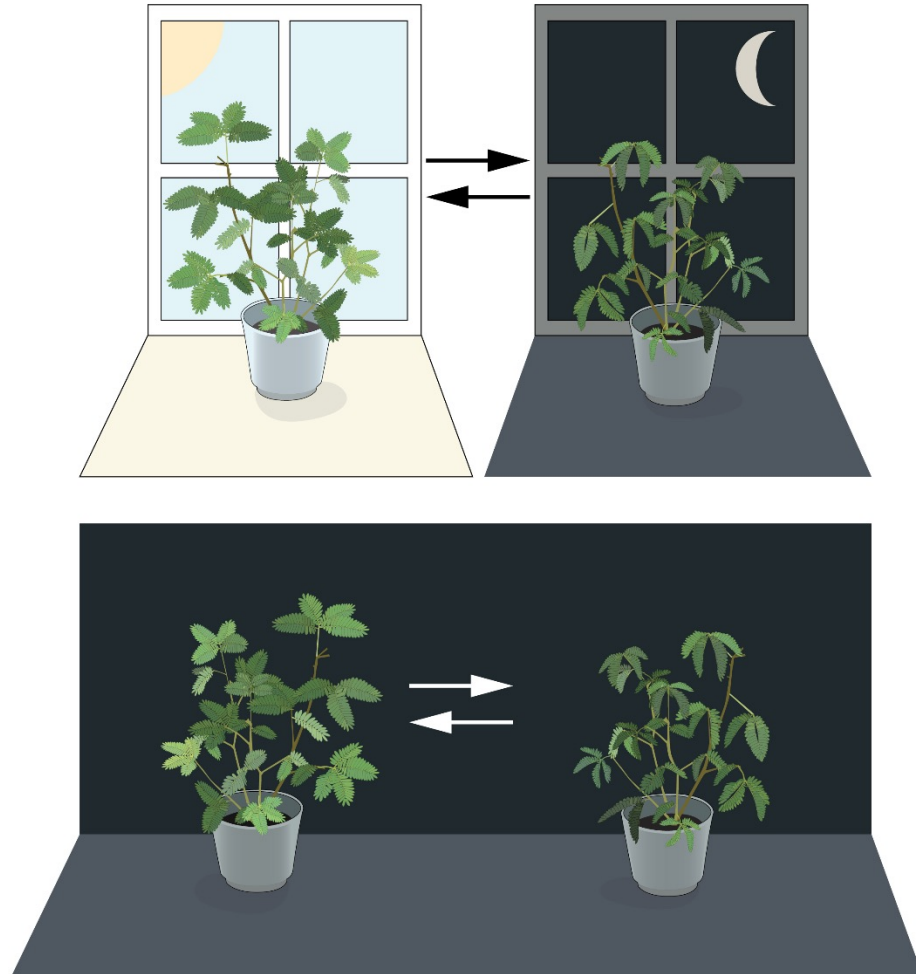


Introduction to translational research

From bench to bedside and back

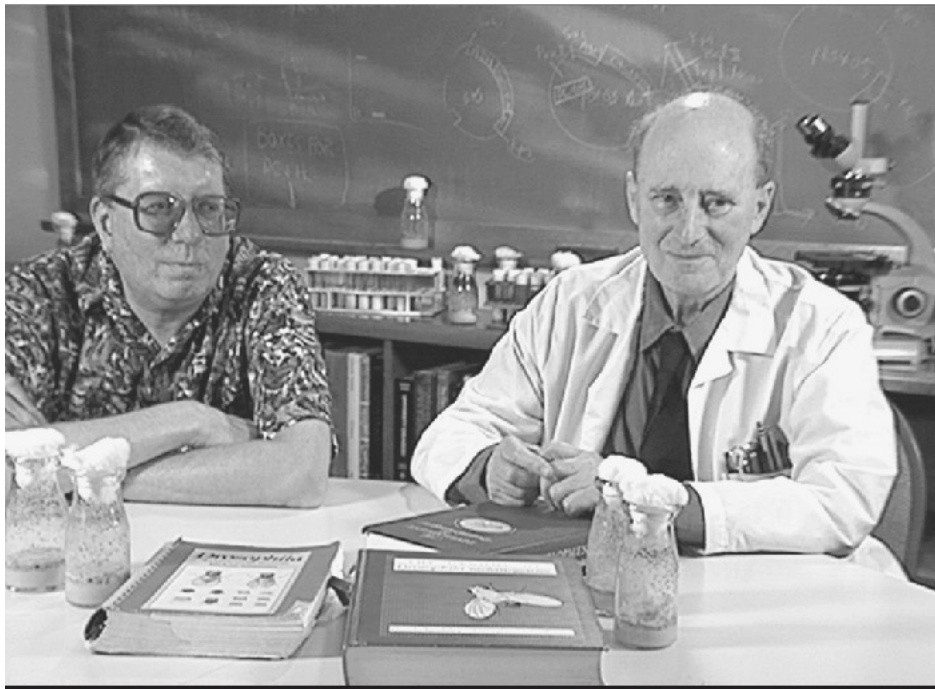
By Solomon Nshemere

First observation of chronobiology in plants



Jacques d'Ortous de Mairan

Discovery of drosophila mutants with disrupted circadian rhythms

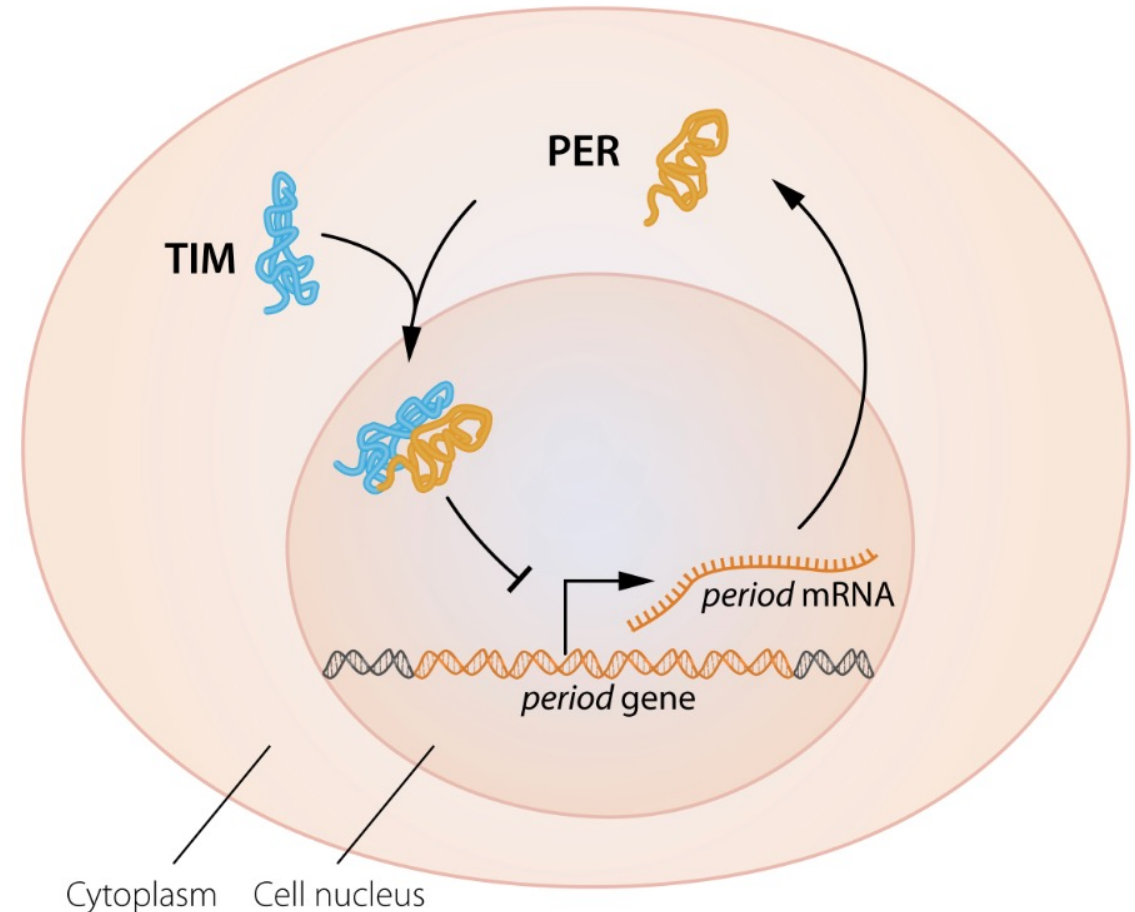


During the 1970's, **Seymour Benzer** and his student **Ronald Konopka** asked whether it would be possible to identify genes that control the circadian rhythm in fruit flies. They demonstrated that mutations in an unknown gene disrupted the circadian clock of flies. They named this gene *period*

Isolation of genes responsible for drosophila circadian biology



In 1984, **Jeffrey Hall** and **Michael Rosbash**, working in close collaboration at Brandeis University in Boston, and **Michael Young** at the Rockefeller University in New York, succeeded in isolating the *period* gene. Jeffrey Hall and Michael Rosbash then went on to discover that PER, the protein encoded by *period*, accumulated during the night and was degraded during the day. Thus, PER protein levels oscillate over a 24-hour cycle, in synchrony with the circadian rhythm



Nobel Prize in Physiology or Medicine 2017



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Jeffrey C. Hall



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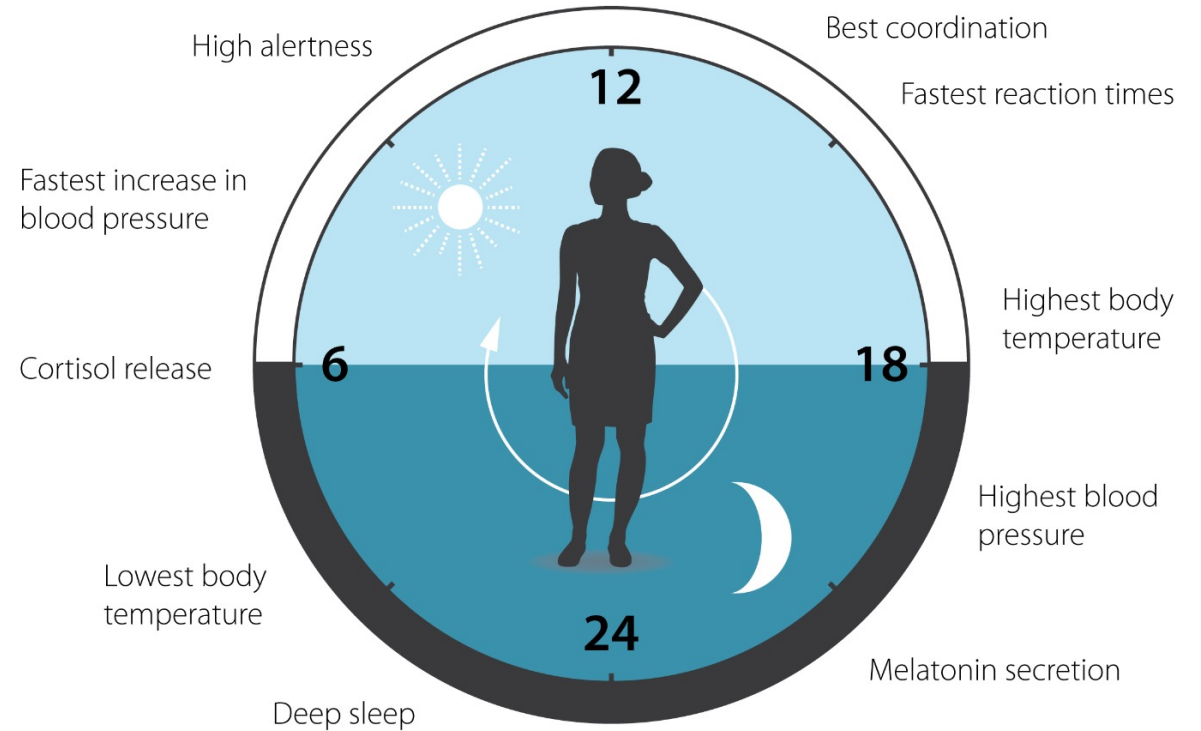
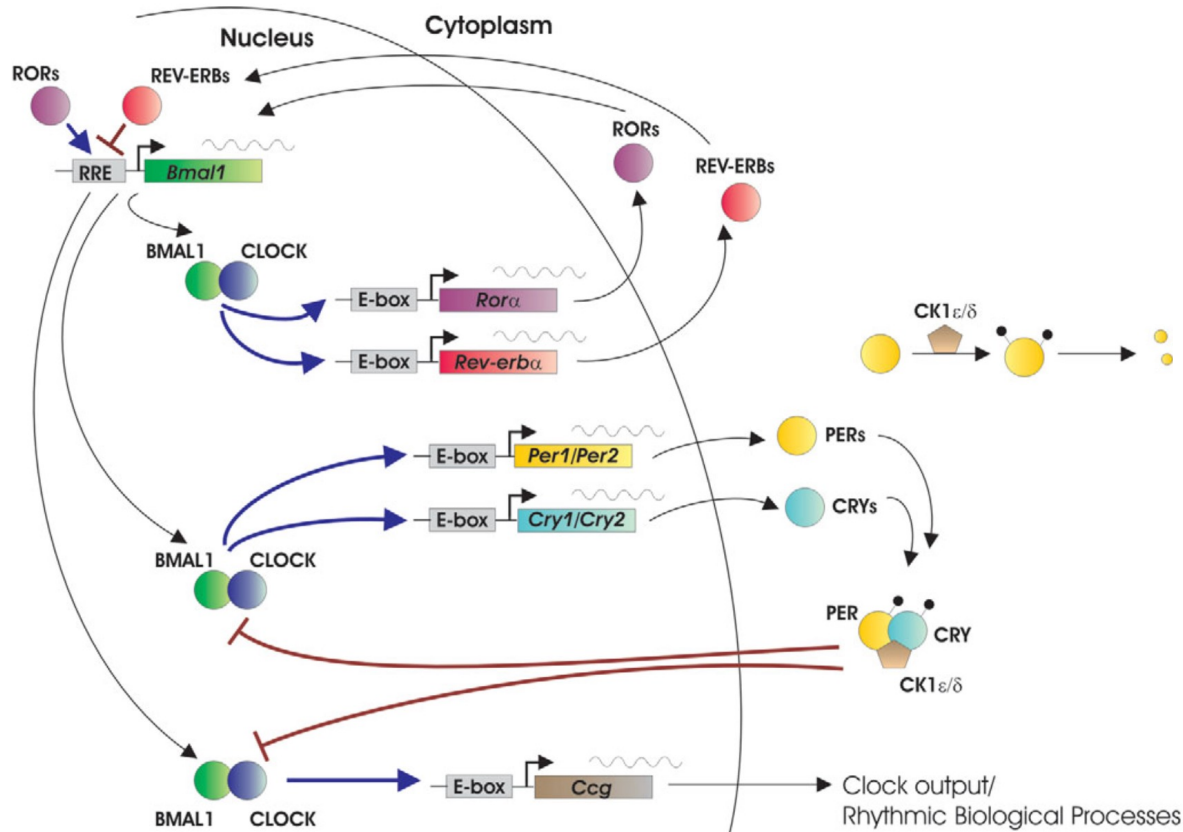
Michael Rosbash



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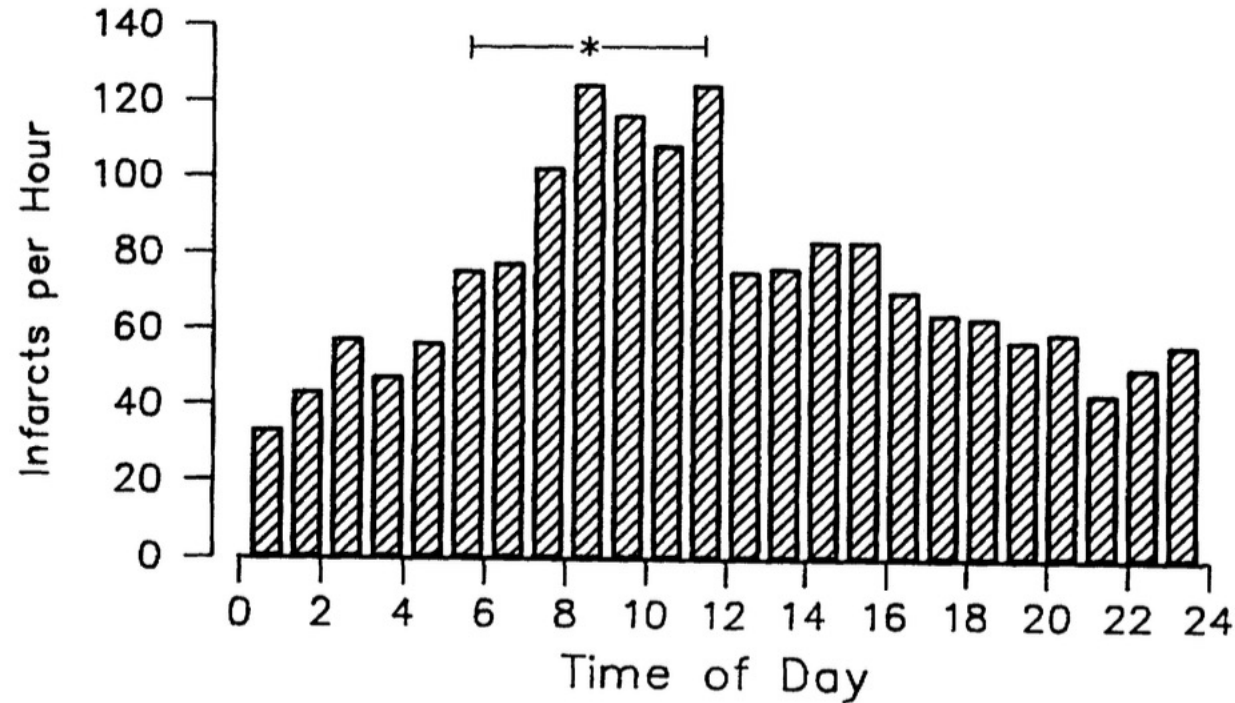
Michael W. Young

The circadian clock in humans has since been elucidated and plays important roles in human physiology and pathophysiology



Myocardial Infarctions show a clear circadian pattern

1, 741 patients



Translating basic research into the clinic

BMAL1–HIF2A heterodimer modulates circadian variations of myocardial injury

<https://doi.org/10.1038/s41586-025-08898-z>

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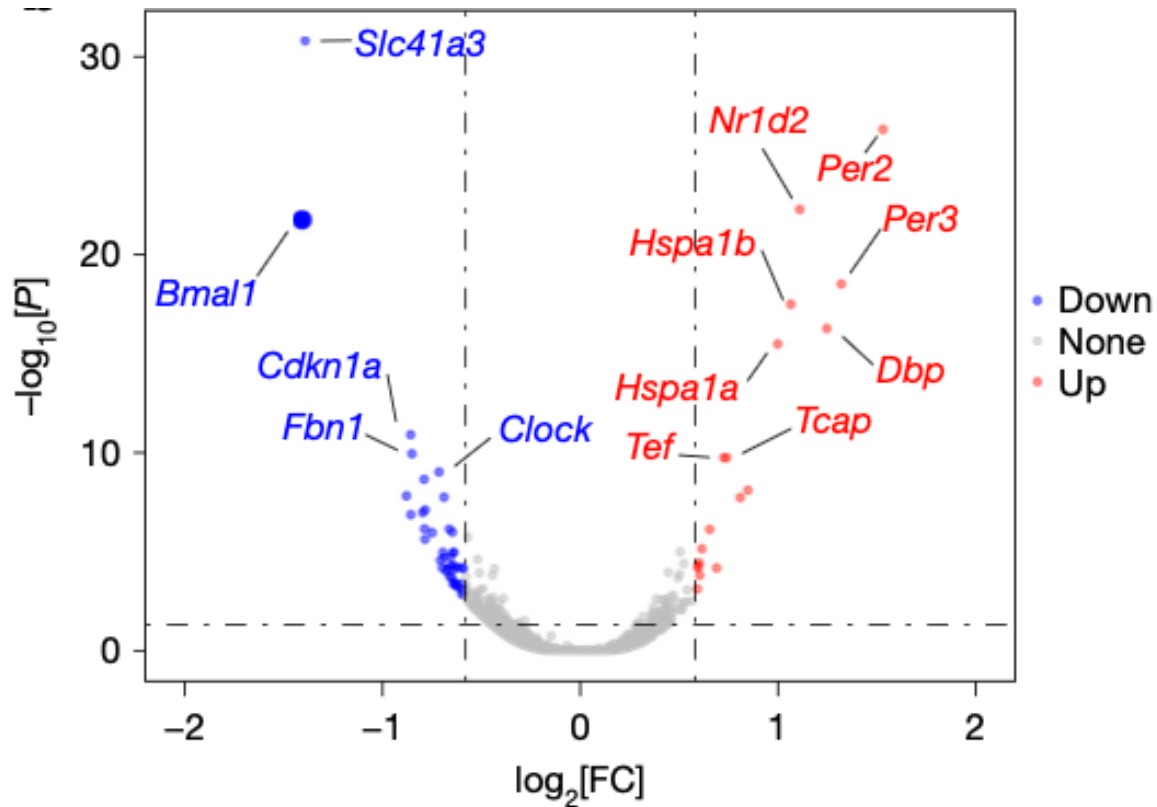
 Check for updates

Wei Ruan^{1,2,18}✉, Tao Li^{3,18}, In Hyuk Bang^{1,18}, Jaewoong Lee^{4,18}, Wankun Deng⁵, Xinxin Ma¹, Cong Luo^{1,2}, Fang Du^{1,6}, Seung-Hee Yoo³, Boyun Kim^{1,7}, Jiwen Li^{1,8}, Xiaoyi Yuan¹, Katherine Figarella¹, Yu A. An¹, Yin-Ying Wang⁵, Yafen Liang^{1,9}, Matthew DeBerge¹, Dongze Zhang¹, Zhen Zhou¹⁰, Yanyu Wang¹, Joshua M. Gorham¹¹, Jonathan G. Seidman¹¹, Christine E. Seidman¹¹, Sary F. Aranki¹², Ragini Nair¹, Lei Li¹³, Jagat Narula¹⁴, Zhongming Zhao⁵, Alemayehu A. Gorfe¹⁵, Jochen D. Muehlschlegel¹⁶, Kuang-Lei Tsai^{1,17}✉ & Holger K. Eltzschig^{1,9}✉

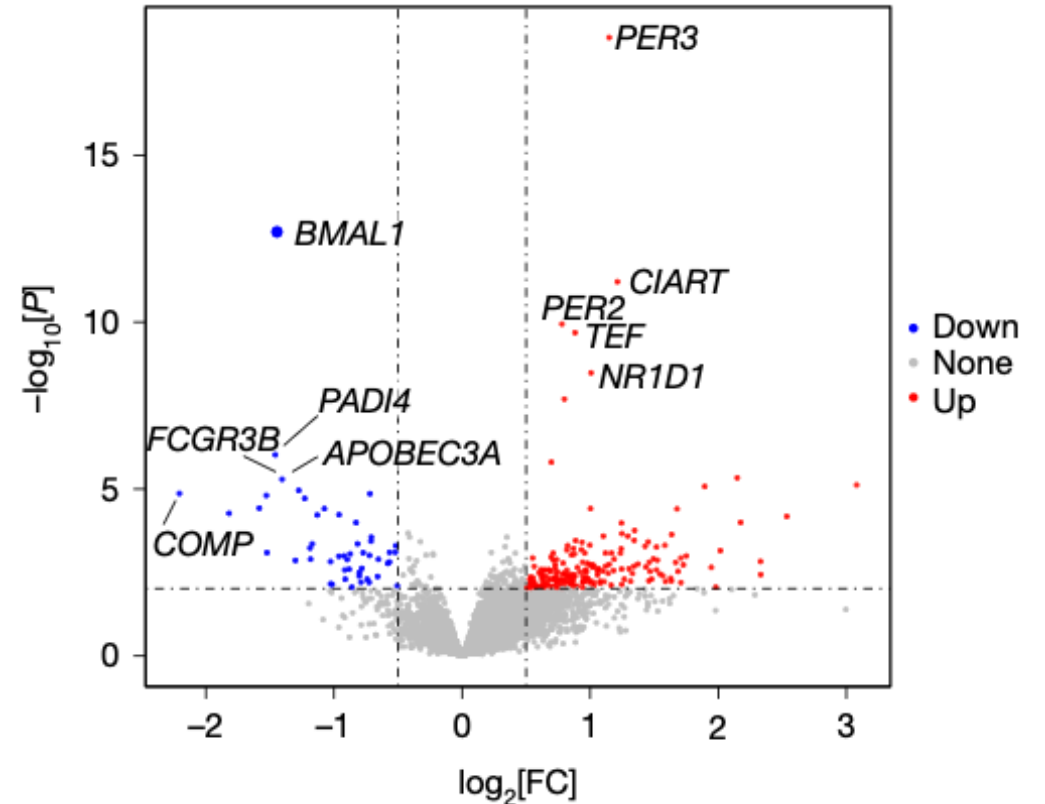
Acute myocardial infarction is a leading cause of morbidity and mortality worldwide¹. Clinical studies have shown that the severity of cardiac injury after myocardial infarction exhibits a circadian pattern, with larger infarcts and poorer outcomes in patients experiencing morning-onset events^{2–7}. However, the molecular mechanisms underlying these diurnal variations remain unclear. **Here we show that the core circadian transcription factor BMAL1^{7–11} regulates circadian-dependent myocardial injury by forming a transcriptionally active heterodimer with a non-canonical partner—hypoxia-inducible factor 2 alpha (HIF2A)^{12–16}—in a diurnal manner.** To substantiate this finding, we determined the cryo-EM structure of the BMAL1–HIF2A–DNA complex, revealing structural rearrangements within BMAL1 that enable cross-talk between circadian rhythms and hypoxia signalling. BMAL1 modulates the circadian hypoxic response by enhancing the transcriptional activity of HIF2A and stabilizing the HIF2A protein. **We further identified amphiregulin (AREG)^{16,17} as a rhythmic target of the BMAL1–HIF2A complex, critical for regulating daytime variations of myocardial injury. Pharmacologically targeting the BMAL1–HIF2A–AREG pathway provides cardioprotection, with maximum efficacy when aligned with the pathway’s circadian phase.** These findings identify a mechanism governing circadian

Ruan, W., Li, T., Bang, I.H. *et al.* BMAL1–HIF2A heterodimer modulates circadian variations of myocardial injury. *Nature* **641**, 1017–1028 (2025)

Mouse models can recapitulate circadian rhythms observed in humans



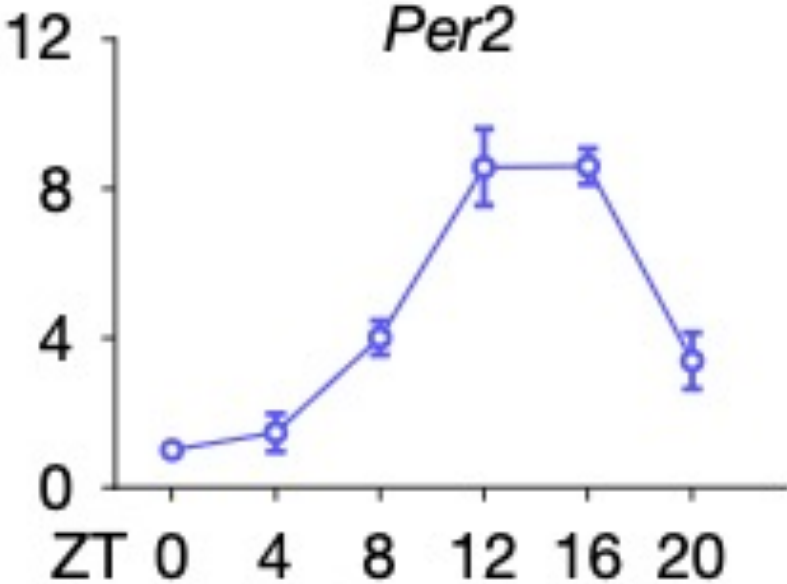
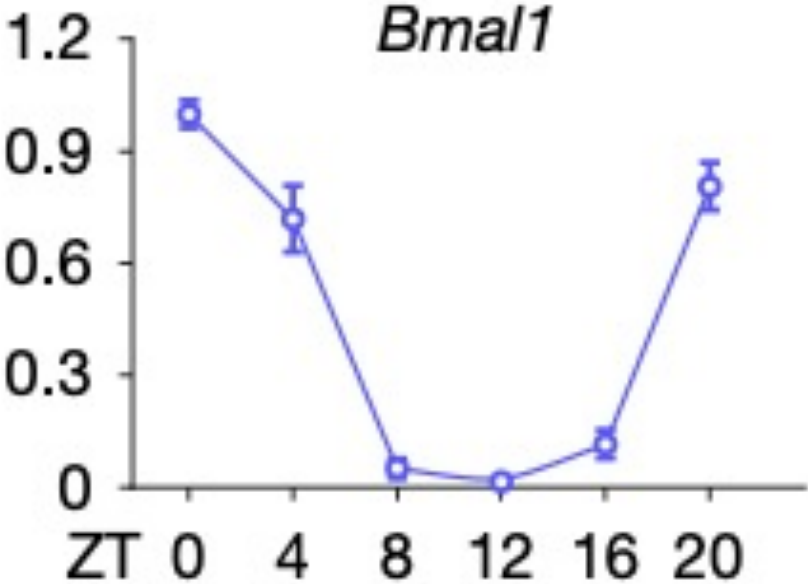
RNA seq of AAR samples taken from mice at ZT8 when MI is least prominent



RNA seq of ischemic patient samples taken in the morning (10:32 am)

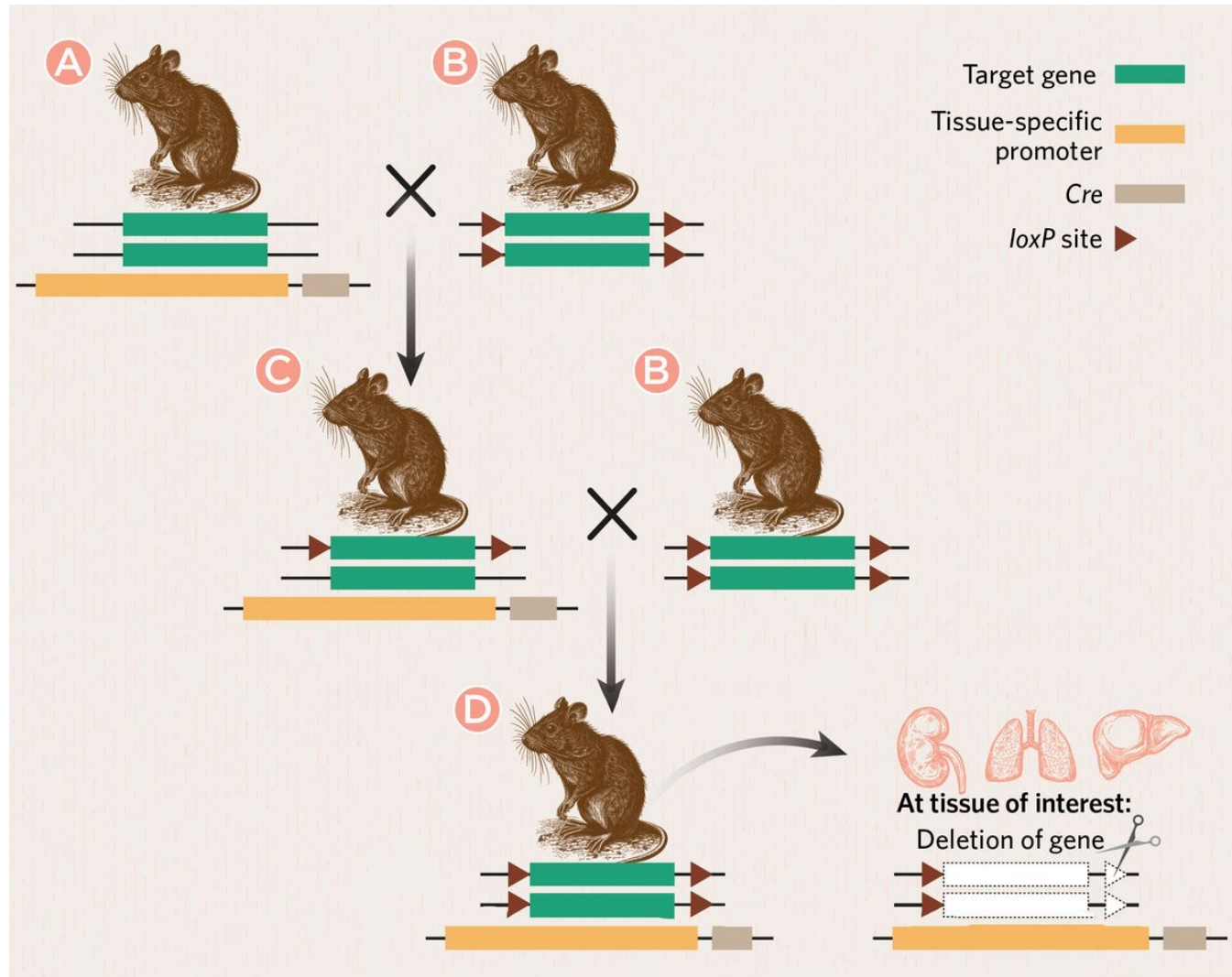
Bmal1 and Per2 show circadian patterns in mice similar to humans

Relative gene expression



Zeitgeber time

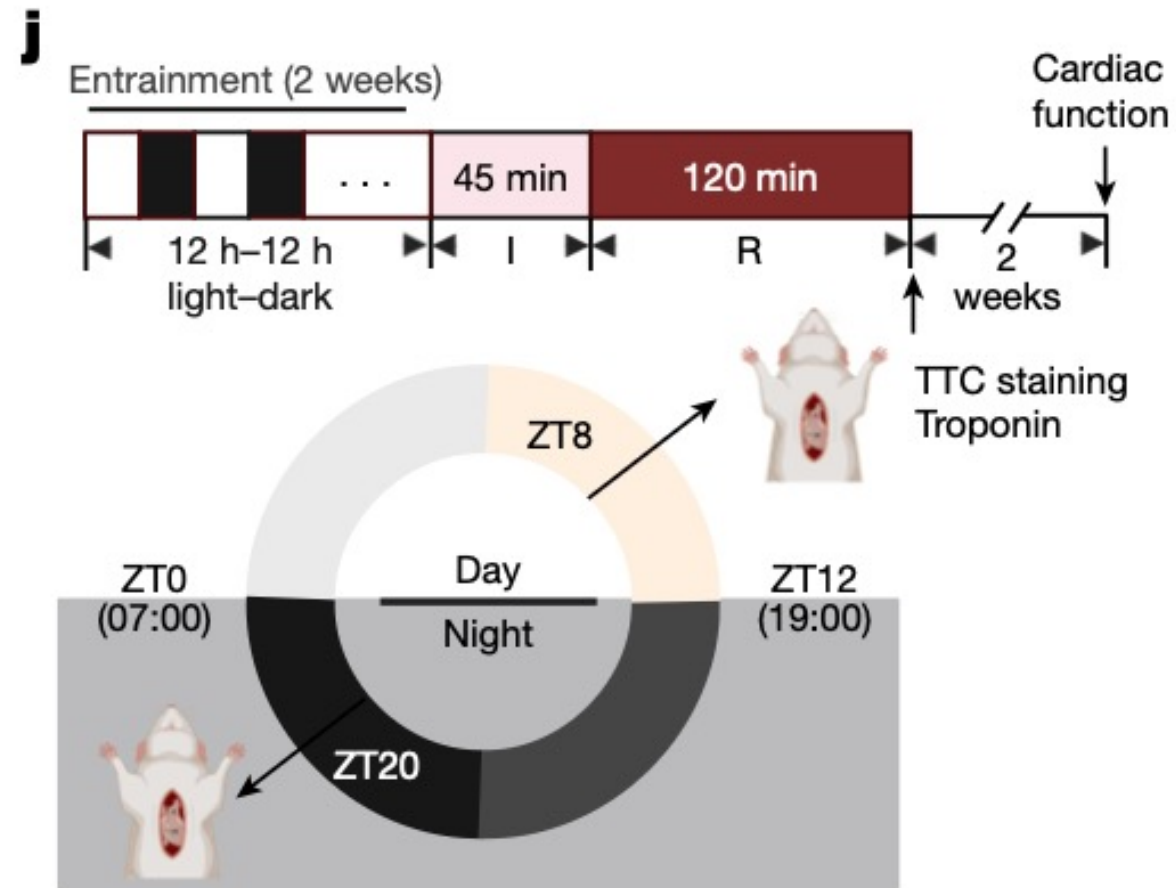
Generation of transgenic mice to investigate Bmal1's functional role in cardiac injury



Bmal1
Myh6
Cre recombinase

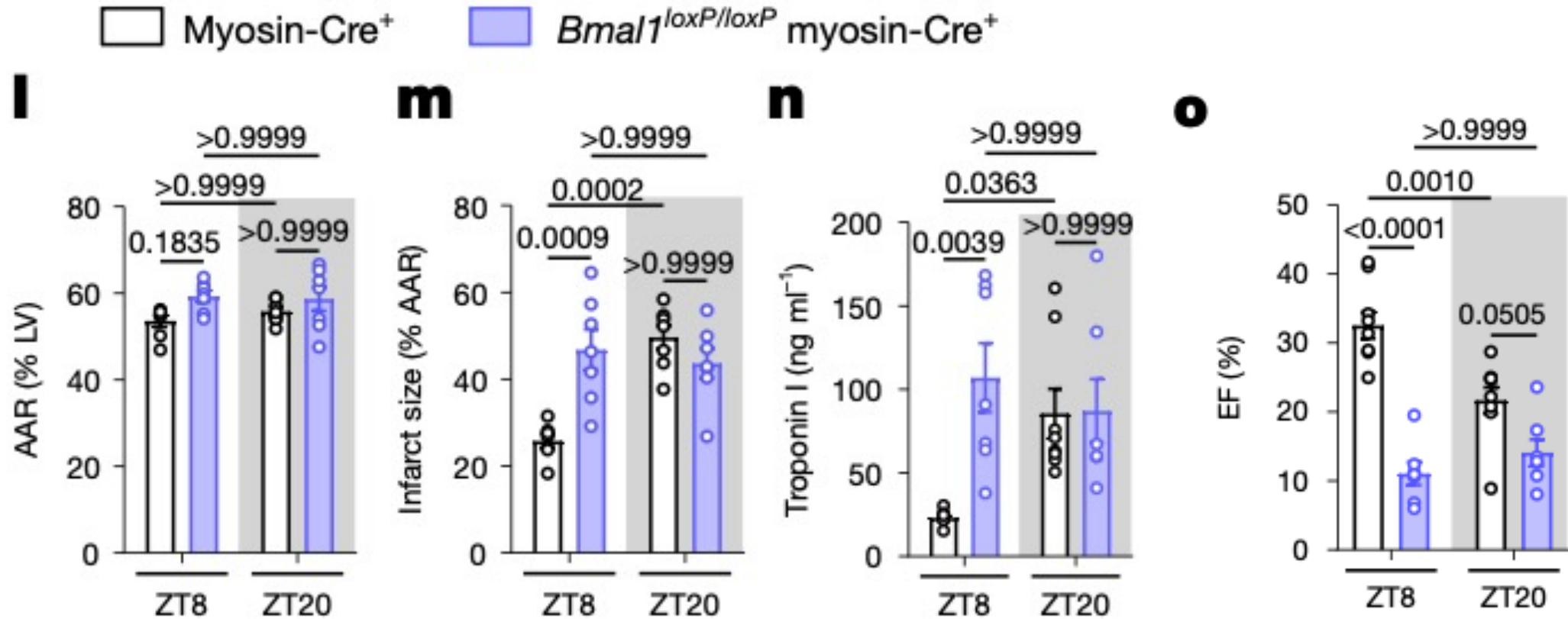
The result is a mouse model in which tamoxifen injection will induce ablation of Bmal1 specifically in cardiomyocytes

Study design



Bmal1 KO mice and Control mice were subjected to MI either at ZT8 or ZT20

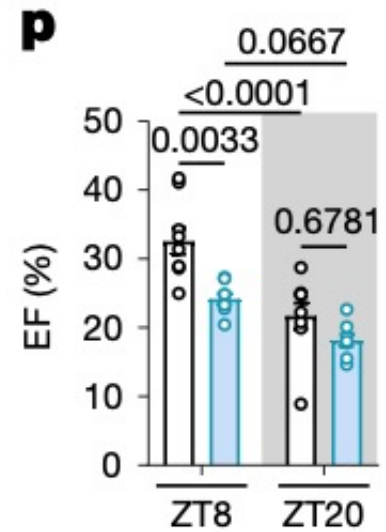
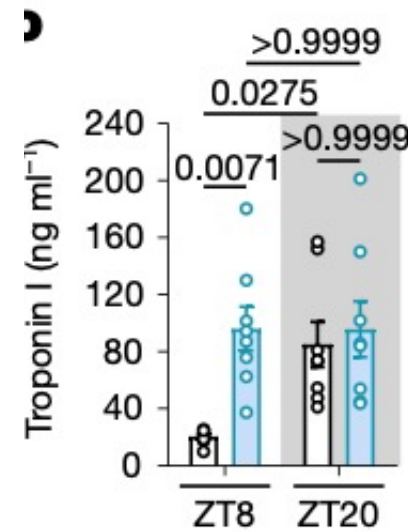
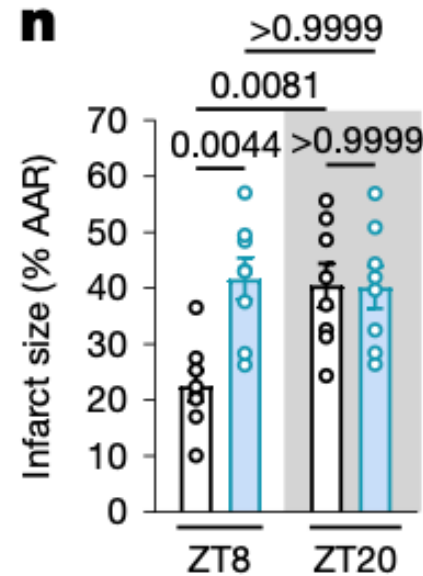
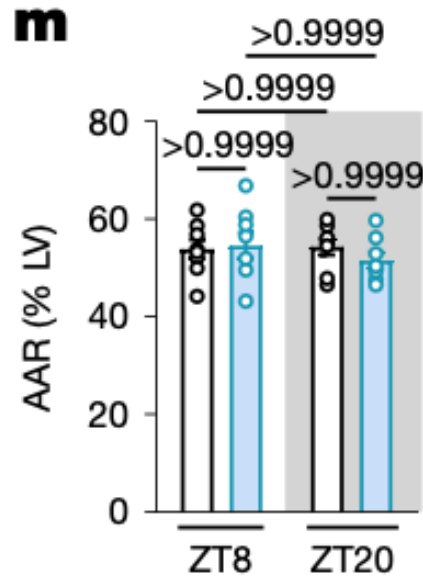
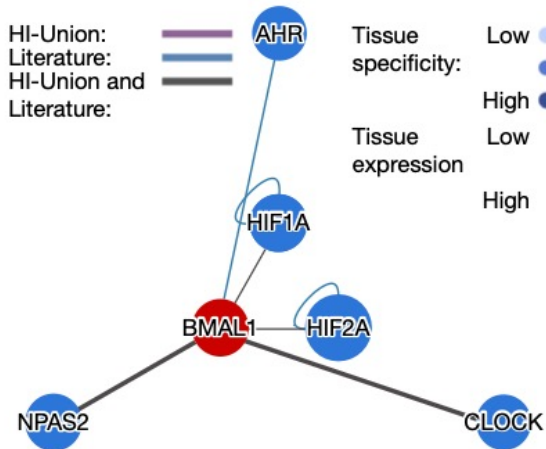
Bmal1 regulates circadian cardiac injury



But how does Bmal1 regulate circadian cardiac injury?

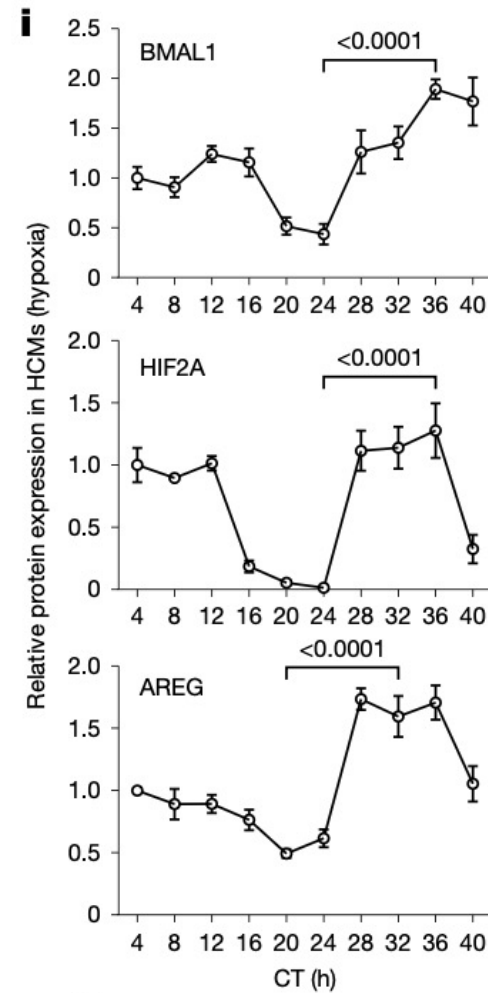
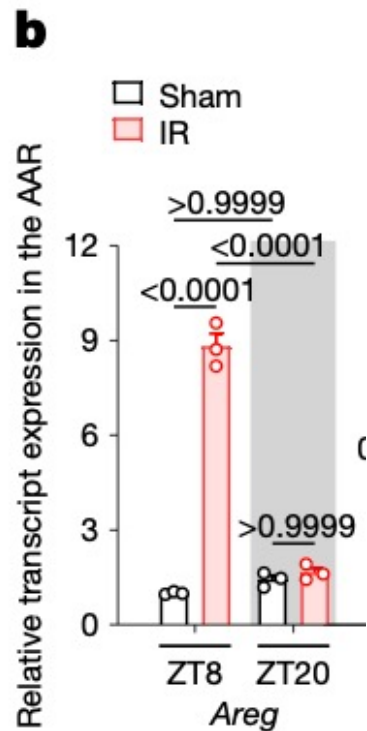
Bmal1 interacts with Hif2a to regulate circadian cardiac injury

Using Cre-Lox strategy, the authors created a mouse model in which **Hif2a is ablated in cardiomyocytes** upon tamoxifen injection

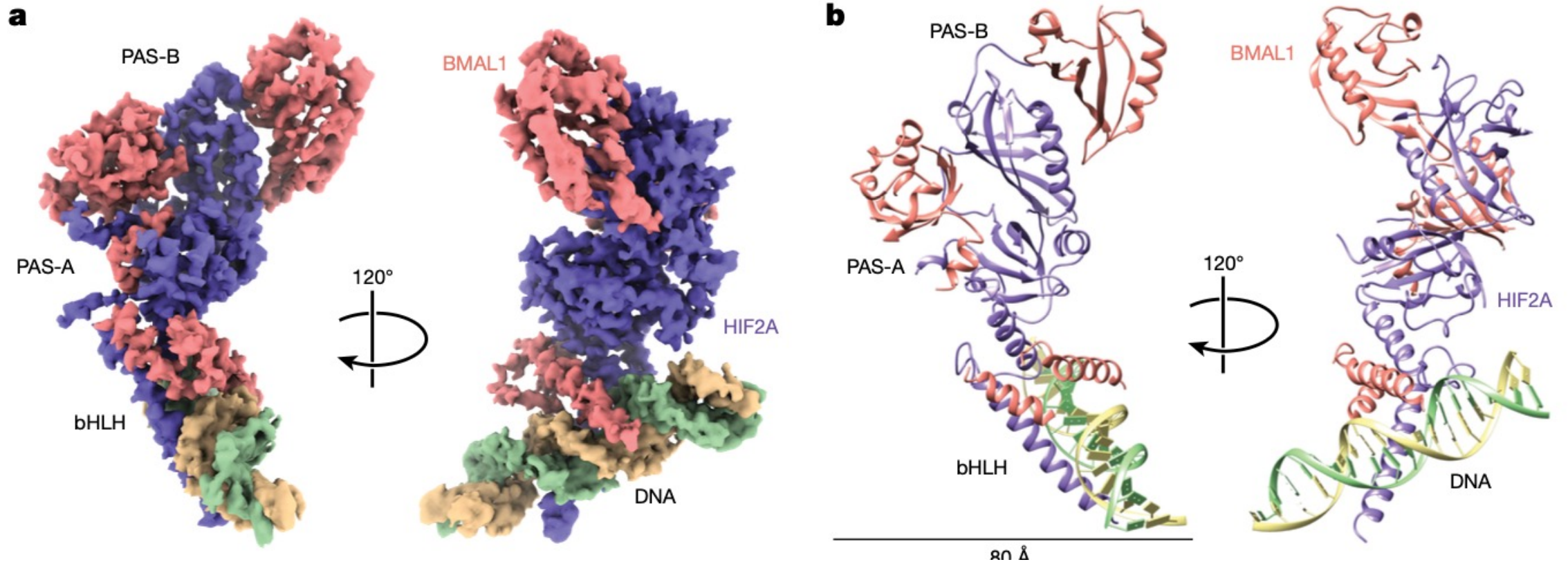


But how does Bmal-Hif2a
complex regulate circadian
cardiac injury?

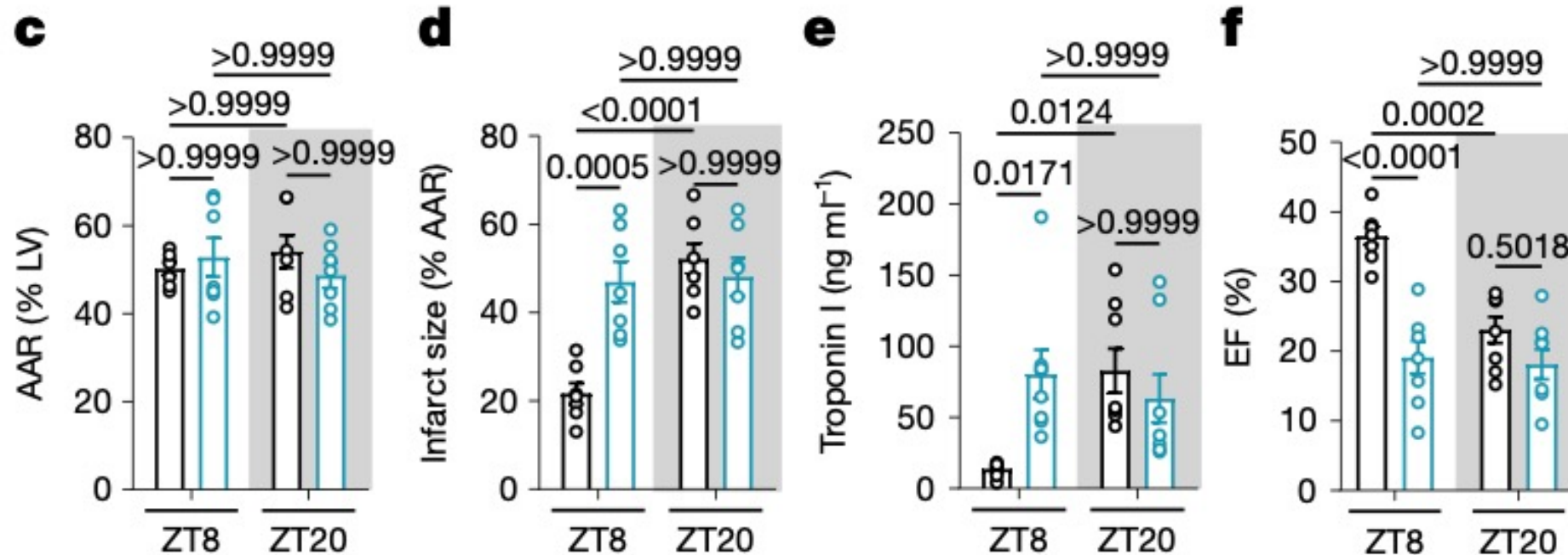
Areg is the most significantly upregulated Hif2a target gene after cardiac injury and shows similar circadian patterns



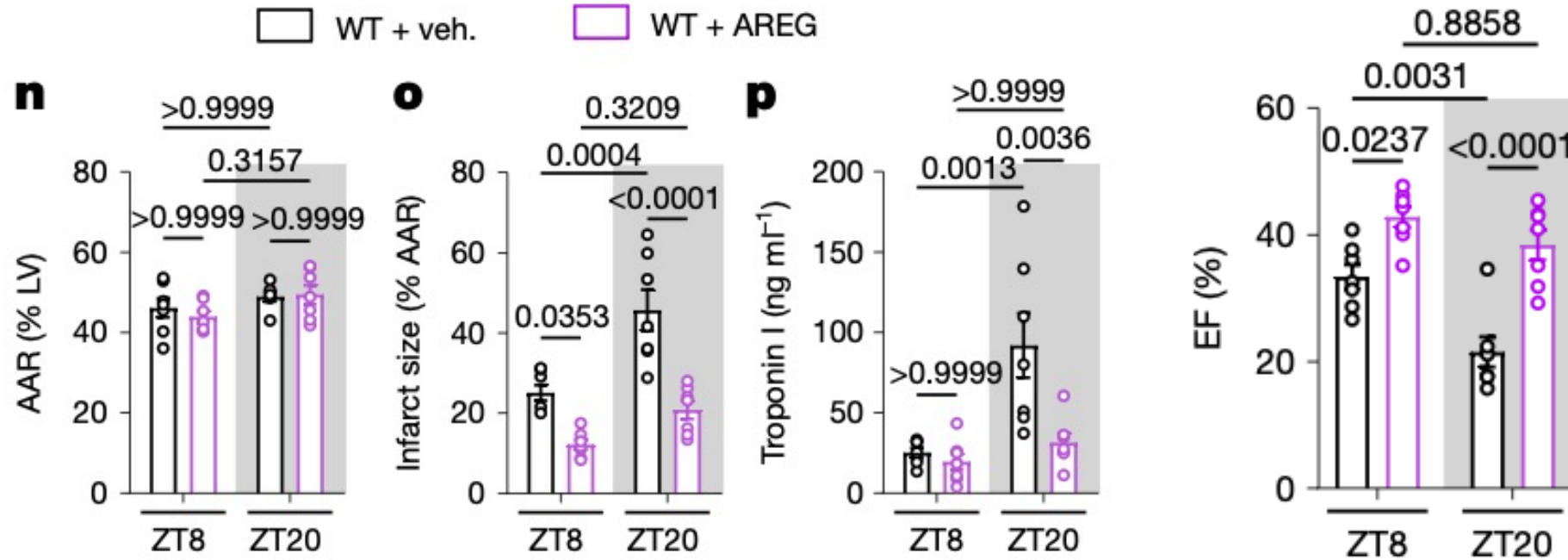
The authors show that Bmal1 dimerizes with Hif2a to drive Areg expression



Areg KO mice exhibit similar characteristics to Bmal1 and Hif2a KO mice



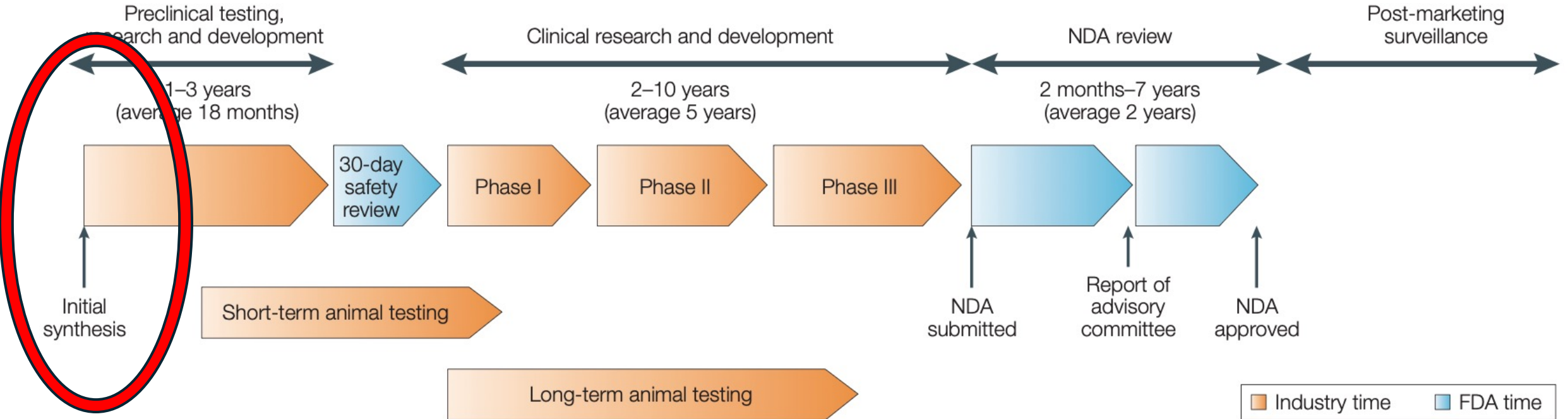
Recombinant Areg as a chronotherapeutic strategy??



Conclusions summary

- The authors elucidate the mechanism by which **Bmal1 regulates cardiac circadian cycles by interacting with Bmal1 to drive expression of Areg**
- The authors elegantly demonstrate that **targeting the Bmal1-Hif2a-Areg axis could be a potential therapeutic avenue** for myocardial injury

So where does this work place us in the drug development timeline?



Thank you for listening