Aficamten, a Selective Small-Molecule Cardiac Myosin Inhibitor for the Potential Treatment of Hypertrophic Cardiomyopathy

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ABSTRACT

Mutations in the cardiac sarcomere increase its contractility and appear to drive the pathogenesis of most forms of hypertrophic cardiomyopathy (HCM). Directly targeting this pathophysiology improves symptoms and exercise capacity in patients with obstructive HCM, as demonstrated by the first-inclass cardiac myosin inhibitor, *mavacamten*. Here we describe the next-in-class selective small molecule inhibitor of cardiac myosin, aficamten (CK-3773274). Aficamten inhibits the myosin ATPase by slowing phosphate release and stabilizing a weak actin-binding state of myosin. Binding to an allosteric site on the myosin catalytic domain distinct from that of *mavacamten*, *aficamten* prevents the conformational changes necessary to enter the strongly actin-bound force-generating state. In single ATP turnover experiments, *aficamten* slowed ATP turnover to rates to <1/3 of the super-relaxed state (SRX) rate. The expected consequence of this mechanism of action is to reduce the number of functional myosin heads that drive sarcomere shortening. *Aficamten* reduced cardiac contractility in cardiac myocytes and in preclinical models of cardiac function. Cardiac myosin inhibition with *aficamten* may offer an alternative approach to treat the underlying hypercontractility of the cardiac sarcomere in HCM.

INTRODUCTION

- Hypercontractility of the cardiac sarcomere appears to underlie pathological hypertrophy and fibrosis in select genetic hypertrophic cardiomyopathies.
- Direct modulation of the sarcomere is an emerging approach to potentially treat conditions with maladaptive changes in cardiac contractility (Malik et al. 2011, Green et al. 2016).
- Clinically, in phase 2 studies, the next-in-class small-molecule inhibitor of cardiac myosin, *aficamten*, has been shown to reduce gradients, improve functional capacity, and lead to improved cardiac remodeling (Maron et al. 2021).
- The objective of this study was to characterize the biochemical mechanism of action of the small molecule *aficamten*, and the ability to modulate cardiac contractility in vitro and in vivo in preclinical models of cardiac function.

METHODS

Preparation of Reagents • Myofibrils were prepared from flash-frozen bovine cardiac, bovine masseter, and rabbit psoas tissue as described in Hwee et al. (2015). Bovine cardiac myosin subfragment-1 (S1) was prepared as described in Malik et al. (2011). Bovine cardiac HMM was prepared based on the method of Rohde et al. (2018). Chicken gizzard myosin and subfragment-1 were prepared as described in Sellers et al. (1981).

ATPase Assays

• Steady-state ATPase activity was measured using a pyruvate kinase and lactate dehydrogenase-coupled enzyme system as described in Hwee et al. (2015) and Malik et al. (2011). Non-myosin ATPase activity was subtracted from cardiac and slow skeletal myofibril assays (where indicated) by subtracting the ATPase activity in the presence of a saturating concentration of the non-selective myosin ll inhibitor (-)-blebbistatin.

Blebbistatin Binding Assays

 Binding of (-)-blebbistatin to bovine cardiac myosin subfragment-1 was measured in a buffer consisting of 12 mM K-PIPES pH 6.8, 2 mM MgCl₂, 1 mM DTT, and 2 mM ADP-vanadate. Fluorescence emission spectra were recorded using a PTI QM-6 (λ_{ex} 426 nm). Compound titrations were performed using a SpectraMax Gemini XS spectrofluorimeter (λ_{ex} 426 nm, λ_{em} 575 nm).

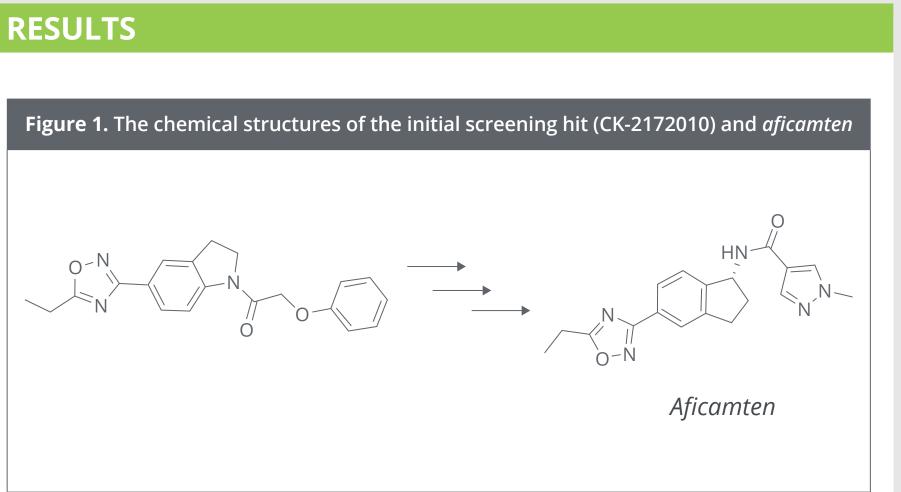
Transient Kinetics

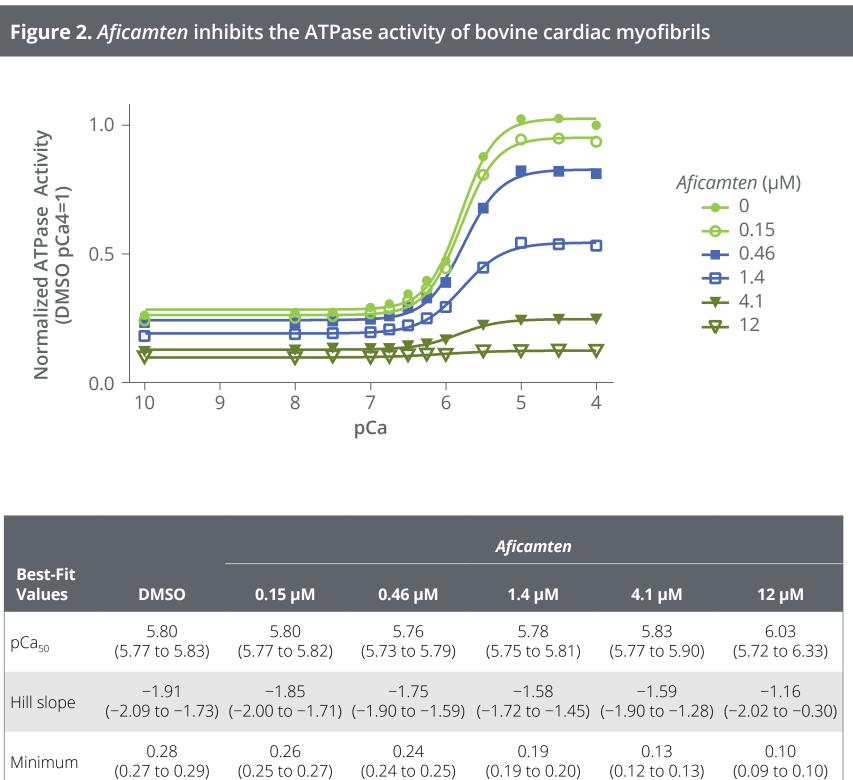
 Transient kinetics were measured at 25°C using an SF-61DX stopped-flow instrument (TgK Scientific) in a buffer consisting of 12 mM K-PIPES pH 6.8, 2 mM MgCl₂, and 1 mM DTT using the methods of De La Cruz and Ostap (2009) and Malik et al. (2011). ATP binding and hydrolysis were monitored by intrinsic tryptophan fluorescence (λ_{ex} 295 nm) using a 320 nm longpass filter. Actinactivated phosphate release was monitored using MDCC-labeled phosphate binding protein (λ_{ex} 434 nm) using a 455 nm longpass filter. Single ATP turnover measurements were performed by mixing mantATP-loaded myosin with 1 mM ATP either in a stopped-flow instrument (Hi-Tech Scientific SF-61 DX2, λ_{ex} 374 nm/ λ_{em} >455 nm) or plate-based fluorescence reader (Molecular Devices SpectraMax Gemini XS, λ_{av} 405 nm/ λ_{am} 455 nm) (Adhikari et al. 2019; Sarkar et al. 2020) in the above buffer supplemented with 60 mM KCl.

Measurement of Cardiomyocyte Contractility and

- **Calcium Transients** • Adult rat ventricular cardiomyocytes were isolated and loaded with Fura-2 as described in Malik et al. (2011). Cardiomyocyte contractility and calcium transients were measured by edgedetection video microscopy and fluorescence photometry (IonOptix, Milton, MA) as described in Malik et al. (2011).
- Rat Echocardiography Assessment
- Adult male Sprague Dawley rats were anesthetized with inhaled isoflurane (1-5%) throughout the echocardiography procedure. Baseline contractility was assessed 1 day prior to *aficamten* treatment. Animals were orally dosed with vehicle (0.5% hydroxypropylmethylcellulose (HPMC)/0.1% Tween-80) or aficamten (0.5, 1, 2, or 4 mg/kg) and measures of left ventricular contractility were assessed 1, 4, 8, and 24 hours post-dose. Using a GE Vivid7 machine, a 10 MHz probe was placed at the level of the papillary muscles and 2D M-mode images of the left ventricle were captured. Images and measurements were obtained in parasternal long axis view. In vivo percent fractional shortening was determined by analysis of the M-mode images using the GE Vivid7 ultrasound software.

Chuang et al. (2021).





Best-Fit Values	DMSO	 0.15 μM	0.46 µl
	BIIISO	0.15 μ	
pCa ₅₀	5.80	5.80	5.76
	(5.77 to 5.83)	(5.77 to 5.82)	(5.73 to 5
Hill slope	-1.91	-1.85	-1.75
	(-2.09 to -1.73)	(-2.00 to -1.71)	(-1.90 to -
Minimum	0.28	0.26	0.24
	(0.27 to 0.29)	(0.25 to 0.27)	(0.24 to C
Maximum	1.03	0.95	0.83
	(1.01 to 1.04)	(0.94 to 0.96)	(0.82 to C
-			

• The ATPase activity of Triton X-100-skinned bovine cardiac myofibrils was measured using a pyruvate kinase/lactate dehydrogenase-coupled assay as described in Hwee et al. (2015). The table shows the results of fitting to a four-parameter dose-response equation along with the 95% confidence interval (CI) of the fitted parameters.

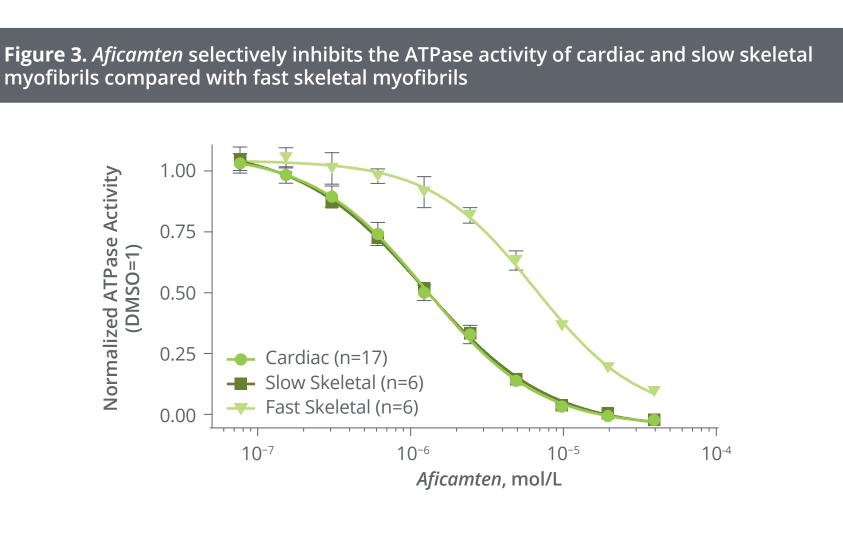
0.54

0.25

0.84) (0.54 to 0.55) (0.24 to 0.25) (0.12 to 0.13)

0.12

Data shown are mean values \pm standard deviation (SD) (n=4)



	Cardiac	Slow Skeletal	Fast Skeletal
IC ₅₀ (μΜ) (95% CI)	1.26	1.23	6.52
	(1.20 to 1.33)	(1.17 to 1.29)	(5.72 to 7.71)
Hill slope	-1.16	-1.06	-1.27
	(-1.24 to -1.10)	(-1.12 to -1.00)	(-1.49 to -1.08)
Тор	1.07	1.10	1.04
	(1.05 to 1.09)	(1.08 to 1.12)	(1.02 to 1.07)
Bottom	-0.04	-0.05	-0.005
	(-0.06 to -0.03)	(-0.07 to -0.04)	(-0.09 to 0.06)

• Dose-response analysis was performed with cardiac (bovine, n=17), slow skeletal (bovine, n=6), and fast skeletal (rabbit, n=6) detergent-skinned myofibrils as described in Hwee et al. (2015). Free Ca²⁺ concentrations were fixed at approximately the pCa₇₅ for each myofibril type. Raw ATPase rates were normalized to reactions containing an equivalent concentration of DMSO, and for cardiac and slow skeletal reactions, ATPase rates in the presence of the nonselective myosin inhibitor blebbistatin were subtracted to eliminate the effects of non-myosin ATPases.

Data shown are mean values ± SD.

