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Time dependent hypotensive effect of aspirin in mice

Lihong Chen^{#1}, Guangrui Yang^{#2,3}, Jiayang Zhang¹, Baoyin Ren², Soonyew Tang³, Xuanwen Li³, and Garret A. FitzGerald^{3,†}

¹Advanced Institute for Medical Sciences, Dalian Medical University, Dalian, 116044, China

²School of Life Science and Biotechnology, Dalian University of Technology, Dalian, 116024, China

³The Institute for Translational Medicine and Therapeutics, Perelman School of Medicine, University of Pennsylvania, Philadelphia, PA 19104, USA

[#] These authors contributed equally to this work.

Abstract

Objective—Evening, but not morning administration of low dose aspirin (ASA) has been reported to lower blood pressure (BP) in hypertensive patients. The present study was designed to determine whether this phenomenon could be replicated in mice, and if so, whether a time-dependent effect of ASA on BP was due to alteration of circadian clock function.

Approach and Results—We recapitulated the protective effect of ASA (50µg/d for 7 days) at Zeitgeber Time (ZT) 0 (active-to-rest transit), but not at ZT12, on a high salt diet-induced increase of blood pressure. However, the time of ASA administration did not influence expression of canonical clock genes or their acetylation. We used mouse Bmal1 and Per2-luciferase reporters expressed in U2OS cells to determine the real-time effect of ASA on circadian function but found that the oscillation of bioluminescence was unaltered. Timing of ASA administration also failed to alter urinary prostaglandin metabolites or catecholamines, or the acetylation of its cyclooxygenase-1 (COX-1) target in platelets.

Conclusions——The time dependent hypotensive effect of aspirin in humans has been recapitulated in hypertensive mice. However, this does not appear to reflect a direct impact of ASA on circadian clocks or on acetylation of platelet COX-1.

Graphical abstract

Disclosures None.

[†]Address for correspondence: Institute for Translational Medicine and Therapeutics, Smilow Center for Translational Research, West Pavilion, 10th Floor, Room 116, 3400 Civic Center Boulevard, Building 421, Philadelphia, PA 19104-5158 Tel: 215-898-1185 | Fax: 215-573-9135. garret@upenn.edu.



The time dependent hypotensive effect of aspirin does not appear to reflect a direct impact on circadian clocks

Introduction

Aspirin (ASA) is most often used at low doses (~100mg/d) in the secondary prevention of myocardial infarction and stroke[•], an effect explained sufficiently by its inhibition of formation of thromboxane (Tx)A₂, the predominant product of cyclooxygenase (COX)-1 in platelets. Higher doses of aspirin are necessary to inhibit COX-2 and thereby relieve pain and inflammation. Deletion of COX-2 accelerates thrombogenesis and elevates blood pressure. Interestingly, knockdown of COX-1, that simulates the pharmacological effect of low dose ASA in mice, not only delays thrombogenesis, but also restrains the elevation of blood pressure caused by COX-2 inhibition.

In humans, several controlled clinical trials have reported that low dose aspirin reduces blood pressure in hypertensive populations, but only when given in the evening, raising the possibility that this effect may contribute to its cardiovascular benefit in hypertensive populations. However, most patients continue to take aspirin in the morning. This may reflect caution emanating from a failure to replicate these findings that mostly derive from the work of one group of investigators and a failure to identify a mechanism that might explain this phenomenon. Recently, administration of low dose (150mg/day) aspirin in the evening was reported to reduce the incidence of pre-eclampsia in a susceptible population, but the study did not include a control group given aspirin in the morning.

Here we wished to determine if we could recapitulate this phenomenon in mice, better to address potential mechanism. Given the broad ranging impact of the molecular clock on drug targets and drug disposition, and the influence of the clock on blood pressure and its regulatory mechanisms, we wondered if this might explain an antihypertensive effect of aspirin conditioned by the time of dosing. As ASA can acetylate proteins beyond platelet COX-1 and because that the function of some core clock proteins can be modulated by acetylation, we hypothesized that a direct impact of ASA on the circadian clock might explain its time-dependent effects on blood pressure.

Materials and Methods

The data that support the findings of this study are available from the corresponding author upon reasonable request.

Experimental design for BP recording

12-week-old male C57BL/6 mice acclimatized to a 12/12-h light/dark cycle for 1 week before radio-telemetry implantation. Although our studies were confined to male mice, the clinical reports of time dependent hypotensive effects of low dose aspirin are evident in both genders. The basal blood pressure (BP) and heart rate (HR) recordings began 7 days after this operation. Mice were subjected to a high salt diet (HSD, 8% NaCl) for 2 weeks to evoke a hypertensive response, better to clarify the phenotype, then divided randomly into two groups and treated with ASA (50µg/d in 100µl of water) by gavage at Zeitgeber Time (ZT) 0 or ZT12 for 7 consecutive days followed by BP recording. Urine was collected for 24 hours using metabolic cages after each BP recording. At the end of the experiment, mice were euthanized by CO_2 inhalation for tissue sample collections (Fig. 1). Samples were stored at $-80^{\circ}C$ until use.

A separate set of HSD and aspirin treated mice were used for analysis of gene expression assay. Tissue samples were collected at ZT8 and ZT20 after 2, 4, or 8-day treatment of aspirin. In addition, another set of mice treated with HSD for 4 weeks was used, but for in vivo experiments only, including measurements of BP, HR, and locomotor activity. All procedures were approved by the University of Pennsylvania IACUC and the Animal Center at Dalian Medical University, Dalian, China.

Telemetry recordings

Under general anesthesia, PA-C10 telemetry probes (Data Sciences, St. Paul, MN) were implanted in the carotid artery as previously described. Telemetry recordings began 7 days after surgery. Continuous BP and HR were monitored in unrestricted animals with the use of the Dataquest IV system (Data Sciences).

Urinary prostaglandin (PG) metabolites and catecholamines

Systemic production of PGD₂, PGE₂, PGI₂, and TxA₂ was determined by mass spectrometric quantitation of their major urinary metabolites: PGD-M, PGE-M, PGI-M, and Tx-M, respectively as previously described. Epinephrine (EPI), norepinephrine (NE), and dopamine (DOP) were measured using Elisa (17-TCTHU-E03.1, Alpco Immunoassays) following the manufacturer's instruction. Data were corrected for urinary creatinine.

Quantitative reverse transcription PCR (qRT-PCR)

Total RNA from aorta, heart, kidney and adrenal gland was extracted using RNeasy kit (QIAGEN) following the manufacturer's instructions. The concentration of RNA was measured by nanodrop (Thermo Scientific). 0.5µg of each sample was applied reverse transcription with reagents obtained from Applied Biosystem. Taqman universal PCR Master Mix and probes (Applied Biosystems) for 18s rRNA, COXs, and core clock genes, Bmal1, Clock, Npas2, Reverba, Per1, Per2, Cry1, Cry2, were used for real-time PCR.

Immunoprecipitation and western blot

MOVAS (a mouse aorta smooth muscle cell line) and SVEC4-10 cells (a mouse vascular endothelial cell line) were obtained from American Type Culture Collection (CRL-2797 and –2181, respectively). Primary culture of mouse aorta smooth muscle cells (VSMC) was described previously. At 80% confluence, the cells, grown in a 12-well plate, were treated with 1mM aspirin for 24 hours, and then lysed in cell extraction buffer (Cat. No. FNN0011, Invitrogen). The cell lysates were applied for immunoprecipitation (IP) using rabbit antibodies (bmal1, sc-48790, Santa Cruz; Ac-Lysine, 9814, Cell Signaling Technology) and Dynabeads Protein G (100.03D, Invitrogen) following manufacturer's instruction. The precipitated proteins were eluted using NuPAGE LDS sample buffer (NP0007, Invitrogen) containing a reducing agent (NP0004, Invitrogen) at 70°C. The eluted samples were then resolved by SDS-PAGE gel, transferred to PVDF membranes (LC2002, Invitrogen) and analyzed by using antibodies against bmal1, Ac-lysine, and p53 (sc-126, Santa Cruz). As negative controls, IP was performed in the absence of antibody or in the presence of rabbit IgG.

Quantification of COX-1 acetylation by aspirin

Mice were treated with 50µg ASA by gavage at ZT0 or ZT12. Blood samples were collected 0.5 or 24 hours after treatment and platelets were isolated for measurement of COX-1 acetylation by isotope dilution mass spectrometry in multiple reaction monitoring (MRM) mode as previously described. Briefly, mouse COX-1 specific sequence (MGAPFSLK) and its acetylated form [MGAPFS(+ace)LK] were measured by their unique transitions on a TSQ Vantage triple stage quadrupole mass spectrometer (Thermo Scientific) interfaced with a Nano-ACQUITY UPLC system (Waters). Quantification information was extracted from the peak areas of the transitions using Xcalibur Quan Browser (Thermo Scientific). COX-1 acetylation was calculated using the ratio of peak area of the acetylated peptide divided by total peak area of non-modified peptide and modified peptide.

Clock gene oscillation by Bioluminescence

A clonal line of human osteosarcoma U2OS cells, stably expressing Bmal1-Luc or Per2-Luc reporters was generously provided by John Hogenesch (Cincinnati Children's Hospital Medical Center). Cells were grown in the DMEM medium supplemented with 10% FBS, 0.3 mg/ml L-glutamine, 100 units/ml penicillin, and 100 g/ml streptomycin. For the luminescence recording, the cells were seeded in 3.5cm dishes in culture medium with addition of 0.1 mM luciferin and 0.1µM Dexamethasone. The luminescence was recorded by an automated robotic system (GNF Systems) equipped with a CCD imager (ViewLux, PerkinElmer).

Statistical Analysis

Statistical analysis was performed using Graph Pad Prism 7. All statistical tests were twosided. Kruskal–Wallis one-way ANOVA and a subsequent Dunn's multiple comparisons test was used when a single variable was compared between two or more groups, two-way ANOVA with Sidak multiple comparisons when two variables were compared between groups, and three-way ANOVA with Tukey's multiple comparisons test when three variables

exist as recommended by the software. The cutoff for significance was P < 0.05 (*). **P <0.01, ***P < 0.001. In all figures with error bars unless specified, the graphs depict means \pm SEM.

Results

Administration of aspirin at the active/inactive transition has a protective effect on HSDinduced increase of blood pressure.

Two weeks on the HSD increased systolic BP (SBP) from 101.2 ± 0.9 to 110.6 ± 1.9 mmHg (P<0.05). Mice were then randomly allocated to aspirin (50µg/d) by gavage once per day for 7 consecutive days at times equivalent to awakening (ZT12) or bedtime (ZT0) while maintained on the HSD. The HSD continued to increase SBP in the group given aspirin at ZT12 (110.5 ± 4.1 vs. 116.1 ± 5.4 mmHg, p=0.09, Fig. 2A) with a difference that was more pronounced during the light phase (104.6 ± 4.1 vs. 112.4 ± 5.1 mmHg, p<0.05, Fig. 2B). However, when we administered aspirin at ZT0, the rise in SBP was absent (110.8 ± 1.2 vs. 108.9 ± 2.9 mmHg). This effect was not accompanied by changes of body weight (Fig 2C). After four weeks on the HSD, ASA treatment at ZT0 lowered SBP in both light and dark phases (Fig. 3), and mean arterial pressure (MAP), HR, and locomotor activity in the light phase, while no obvious change was found in those dosed at ZT12 (table 1). Thus, we recapitulated the time dependent hypotensive effect of aspirin in HSD treated mice.

No time dependent effect of aspirin on the expression of circadian genes in mouse heart, aorta, kidney or adrenal gland.

At the end of the in vivo experiments, we collected several organs relevant to BP regulation including heart, aorta, kidney and adrenal gland at ZT8 and ZT20 when both Bmal1 and its reciprocal core clock partner, Rev-erba, show differential gene expression. We found no evidence that time of aspirin dosing differentially impacted expression of the canonical clock genes, Bmal1, Clock, Npas2, Rev-erba, Per1, Per2, Cry1, or Cry2 at either ZT8 or ZT20 at the end of the study (Fig. 4). To address the possibility that there might have been a reversible impact on clock genes at an earlier stage in the dosing period, we performed another experiment in which mice were sacrificed for tissue collection at multiple time points. As shown in Figure I (in the online-only Data Supplement), among all comparisons between ASA-ZT0 and ASA-ZT12, only Per2 in the heart collected on day-4 showed a statistical difference, noted at ZT8.

Aspirin failed to influence directly the molecular clock.

We found that P53 was acetylated or its acetylation was increased by ASA in MOVAS and SVEC4-10 cells. Acetylation of Bmal1 was also detected, however, the pattern was not affected by ASA (Fig. 5A). None of the other clock proteins including Clock, Per1, Per2, Cry1 and Cry2 were found acetylated by ASA treatment. Acetylation of platelet COX-1, the mechanism by which aspirin prevents thrombosis, was unaltered by time of drug administration (Fig. 5B). Using Bmal1-Luc and Per2-Luc reporter expressing U2OS cell lines, ASA (0.1mM, 0.5mM, and 2mM), failed to alter the period, amplitude or phases of the oscillating bioluminescence rhythms (Fig. 6).

No time dependent effect of aspirin on urinary prostaglandin metabolites or catecholamines.

Aspirin administered at both ZT0 and ZT12 decreased prostaglandin biosynthesis as reflected by the urinary metabolites, PGD-M, PGE-M, PGI-M, and Tx-M. However, this impact was unaltered by the time of dosing (Fig. 7A). There was also no time dependent difference in EPI or NE when measured at the end of the study. DOP was lower on aspirin, but there was no difference between ASA-ZT0 and ASA-ZT12 (Fig. 7B).

Discussion

It has long been known that blood pressure (BP) and heart rate (HR) are subject to circadian rhythmicity; indeed, the failure of blood pressure to fall at night ("non-dipping") has been associated with poor cardiovascular clinical outcomes. These time-dependent effects are not just consequences of the sleep/wakefulness cycle, but are believed to be attributable to intrinsic properties of the cardiovascular system that fluctuate under the control of the molecular clock during the course of the day¹. During the last two decades, Hermida and colleagues have reported several randomized trials in hypertensive populations in which low dose aspirin (100mg) administered at night but not in the morning lowered BP in both genders^{1, –}. Recently, in a multicenter, double-blind, placebo-controlled trial, Rolnik et al found that treatment with 150mg aspirin at night in women at risk reduced the incidence of preterm preeclampsia, however they did not control for time of drug administration. Despite these observations, there have been reasons for skepticism. Firstly, some groups have failed to reproduce the fundamental finding¹. Secondly, there is no mechanistic explanation for the phenomenon.

In the current study, we first recapitulated the time-dependent effect of aspirin on human blood pressure in mice. As in humans, ASA lowered blood pressure in hypertensive mice when administered at the transition between activity and rest. Ironically, replication of this phenotype in a model system lends credence to the veracity of the phenomenon in humans. Given the prevalence of hypertension and the effectiveness and commonality of aspirin consumption for the secondary prevention of cardiovascular disease, a population-based shift to evening dosing might have a considerable effect on cardiovascular outcomes. Interestingly, there was no apparent baroreflex mediated increase in heart rate coincident with the ASA mediated time dependent reduction in BP · . Here both HR and activity were decreased by aspirin treatment at ZTO and this may have contributed to the BP lowering effect in mice. Here we saw no impact of ASA on urinary EPI and NE. Although urinary DOP was reduced on ASA, there was no time dependent effect apparent. In future studies we intend to elucidate further the potential contribution of autonomic function as well as addressing other signaling pathways that may underlie these observations.

Indeed, this model affords a more tractable opportunity to address mechanism. The current studies suggest that it does not involve a direct effect of aspirin on the core molecular clock. The time of ASA administration does not obviously influence expression of a variety of clock genes in heart, aorta, kidney or adrenal gland, tissues of direct relevance to blood pressure regulation and cardiovascular homeostasis. Aspirin prevents thrombotic events by acetylating platelet COX-1 but is recognized potentially to acetylate additional proteins.

Although some clock proteins can be modulated by acetylation, ASA did not alter the acetylation patterns of Bmal1 in cultured mouse vascular smooth muscle cells or endothelial cells in vitro. None of the other clock proteins that we studied, including Clock, Per1, Per2, Cry1 and Cry2 were acetylated by ASA treatment. Besides, platelet COX-1 acetylation did not differ at 0.5 or 24 hours after ASA treatment at ZT0 or ZT12. Furthermore, ASA did not alter the oscillation of bioluminescence in U2OS cells expressing Bmal1 and Per2-luciferase reporters. Finally, although rhythmic histone acetylation underlies transcription in the mammalian circadian clock, it is not acetylated by aspirin. These results fail to support the hypothesis that the time dependent effect of ASA is attributable to a direct action on circadian clock.

Aspirin prevents thrombotic events like myocardial infarction by cumulatively acetylating platelet COX-1 dependent TxA₂ formation when administered at low doses that have minimal impact on COX-2 derived prostaglandins like PGI₂. Here the biochemical profile of the dose of aspirin we selected (suppression of urinary metabolites of PGs of the E, D and I series along with Tx) more closely resembles that of anti-inflammatory doses of aspirin in humans. None of the trials of time dependent aspirin administration in humans have used such higher doses although our recapitulation of the functional phenotype (and the failure to implicate platelet COX-1) suggests that the phenomenon may be independent of aspirin dose.

Our findings appear to contrast with the attenuation of the fall in blood pressure during the rest phase in mice globally deficient in COX-1. This may reflect differences in genetic background and/or variability in tissue distribution of aspirin as well as oscillation in its targets that at this dose would involve both COX-1 and COX-2.

In summary, the time dependent impact of aspirin on blood pressure in hypertensive patients was recapitulated in mice made hypertensive by a HSD. This does not result from a direct effect of aspirin on core clock gene expression or oscillation. Recapitulation of this time dependent effect in mice lends weight to the clinical observation and the argument to advise evening dosing with low dose aspirin by hypertensive patients who take it for cardio-protection.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Nonstandard Abbreviations and Acronyms

ASA	aspirin		
BP	blood pressure		
COX	cyclooxygenase		
DOP	dopamine		
EPI	epinephrine		
HR	heart rate		
HSD	high salt diet		
NE	norepinephrine		
PG	prostaglandin		
ZT	Zeitgeber Time		

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- Time dependent hypotensive effect of aspirin in hypertensive patients was recapitulated in mice.
- This effect does not result from a direct impact of aspirin on core clock genes.
- This effect does not reflect a direct impact of aspirin on acetylation of COX-1 or excretion of urinary prostaglandins and catecholamines.

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Figure 1.

Study design for BP recording. Basal BP of 14-week-old mice was recorded using radiotelemetry followed by 2-week HSD treatment. BP was recorded after HSD and mice were then treated with ASA at ZT0 or ZT12 for 7 consecutive days followed by BP recording. Urine was collected after each BP recording. At the end of the experiment, mice were euthanized for tissue collection.

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Figure 2.

A, ASA ($50\mu g/d$ for 7 days) administered at ZTO, but not ZT12, prevented a further increase of systolic blood pressure induced by 2-w HSD (2-way ANOVA; *, p<0.05; **; p<0.01; n=6-7). B, The increase of SBP in ASA-ZT12 group was more pronounced during the light phase. (2-way ANOVA; *, p<0.05; n=6-7). C, There is no statistical difference in body weight between the ASA-ZT0 and ASA-ZT12 dosing groups (2-way ANOVA; n=10).

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Figure 3.

A, ASA (50µg/d for 7 days) administered at ZT0, but not ZT12, lowers high blood pressure induced by 4-w HSD in both light and dark (grey background) phases (2-way ANOVA; *, p<0.05; ns, no statistical difference; n=7-8). B, SBP in A depicted in two consecutive days showed that ASA administered at ZT0 lowers BP at multiple time points across both light and dark phases. (2-way ANOVA; *, p<0.05; n=7-8)

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Figure 4.

mRNA levels of core clock genes in the kidney, heart, aorta, and adrenal gland were revealed by qRT-PCR. All data were normalized to 18s rRNA. There is no obvious difference between ASA-ZT0 and ASA-ZT12 groups in any examined tissues collected either at ZT8 or ZT20 (2-way ANOVA; n=5–6).

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Figure 5.

A, Acetylation pattern of P53 (as a positive control) and Bmal1 in ASA treated blood vessel cells was sought using immunoprecipitation (IP) and Western blotting (WB). MW: molecular weight; B, Mouse platelet COX-1 acetylation did not differ at 0.5 or 24 hours after ASA treatment at ZT0 or ZT12 (2-way ANOVA; ns, no statistical difference; n=4).

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Figure 6.

ASA at multiple concentrations did not alter the oscillation of circadian clocks in Bmal1 (A) or Per2 (B)-luciferase reporter expressed U2OS cells. Similar results were obtained from an independent replication of experiment (1-way ANOVA; n=4).

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Figure 7.

Urinary PG metabolites (A) and catecholamines (B) were measured using mass spectrometry and Elisa, respectively, (2-way ANOVA; *, p<0.05; **; p<0.01; ***, p<0.001; n=9–11). Data were normalized to creatinine.

Table 1.

BP, HR, pulse pressure (PP) and activity parameter averages (±SEM) as measured via radiotelemetry on mice fed with HSD for 4-w before ASA treatment. MAP: mean arterial pressure, DBP: diastolic blood pressure. 2-way ANOVA;

	ASA-ZT0		ASA-ZT12	
	Day	Night	Day	Night
Before ASA treatment				
SBP (mmHg)	112.6±1.52	124.8 ± 2.54	115.1±2.41	125.7±2.96
MAP (mmHg)	99.1±1.13	110.7 ± 2.17	101.3±2.14	111.3±2.55
DBP (mmHg)	85.1±1.23	96.2±2.12	87.4±2.61	96.6±2.62
PP (mmHg)	27.4±1.39	28.5±1.54	28.0±1.77	29.0±1.75
HR (bpm)	470.7±4.70	505.5 ± 4.82	463.9±9.88	531.0±9.94
Activity (arbitrary counts)	3.0±0.40	9.1±2.25	2.7±0.57	9.3±1.79
After ASA treatment				
SBP (mmHg)	108.3±2.24 ^{**}	120.0±2.88 [*]	115.0±2.24	127.6±3.26
MAP (mmHg)	96.3±1.38 ^{**}	107.9±1.92	100.7±2.06	112.9±2.79
DBP (mmHg)	84.2±0.75	95.6±1.31	85.9±2.49	97.8±2.74
PP (mmHg)	24.1±1.83	24.4±2.04	29.2±1.57	29.9±1.68
HR (bpm)	446.0±7.74 ^{**}	503.5±10.64	487.3±8.47	535.4±8.10
Activity (arbitrary counts)	1.7±0.28 ^{**}	6.4±1.03	2.1±0.37	6.8±0.72

** P<0.01,

^{*}P<0.05, after ASA treatment vs. before ASA treatment.