Oxygen and carbon dioxide rhythms are circadian clock-controlled and differentially directed by behavioral signals

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Summary

Daily rhythms in animal physiology are driven by endogenous circadian clocks in part through rest-activity and feeding-fasting cycles. Here, we examined principles that govern daily respiration. We monitored oxygen consumption and carbon dioxide release, as well as tissue oxygenation in freely moving animals to specifically dissect the role of circadian clocks and feeding time on daily respiration. We found that daily rhythms in oxygen and carbon dioxide are clock-controlled and that time-restricted feeding restores their rhythmicity in clock-deficient mice. Remarkably, day-time feeding dissociated oxygen rhythms from carbon dioxide oscillations, whereby oxygen followed activity and carbon dioxide was shifted and aligned with food intake. In addition, changes in carbon dioxide levels altered clock gene expression and phase-shifted the clock. Collectively, our findings indicate that oxygen and carbon dioxide rhythms are clock-controlled and feeding-regulated, and support a potential role for carbon dioxide in phase resetting of peripheral clocks upon feeding.

Introduction

The prevalence of daily rhythms in almost every aspect of biology in mammals has been subject for intensive research in the past years. Among others, these include characterization of daily oscillations in gene expression, protein and metabolite accumulation, as well as overt rhythms in physiology (e.g. core body temperature, heart rate, blood pressure) and behavior (e.g. activity, feeding), (Asher and Sassone-Corsi, 2015; Bass and Lazar, 2016; Curtis and Fitzgerald, 2006; Panda, 2016). These rhythms are governed by endogenous cell autonomous oscillators in conjunction with external inputs such as light-dark cycles that modulate rest-activity and consequently feeding behavior (Dibner et al., 2010; Partch et al., 2014; Takahashi, 2017). The intricate interplay between endogenous rhythms and environmental cues in control of various rhythmic outputs plays a critical role in animal homeostasis. The dissection between these components in the context of whole animal physiology, which readily responds to external inputs and relies as well on endogenous rhythms, is of great interest.

A vital element in mammals' physiology is respiration, namely the exchange of oxygen from the outside environment into cells within tissues in return for carbon dioxide transport in the opposite direction. In cells, the mitochondria utilize nutrients such as carbohydrates and lipids and convert them to energy coins upon oxygen consumption and carbon dioxide release. Oxygen is primarily consumed in mitochondria through oxidative phosphorylation, whereas carbon dioxide is the product of the citric acid cycle (Akram, 2014; Wilson et al., 1988).

This fundamental process exhibits daily rhythms as pulmonary ventilation normally rises during the active phase and drop during the resting phase (Mortola, 2004). Consequently, oxygen consumption rate, as well as blood and tissue oxygenation in mammals exhibit daily oscillations (Adamovich et al., 2017; Emans et al., 2017). We recently reported that oxygen rhythms within the physiological range synchronize circadian clocks in HIF1α-dependent manner and that modulation of ambient oxygen levels accelerate jet-lag adaption (Adamovich et al., 2017). Bass

and colleagues further demonstrated that HIF1α regulate the circadian period and that circadian clock-disruption impairs oxygen sensing through HIF1α (Peek et al., 2017). Furthermore, an interplay between the DNA-binding activity of HIF1α and the core clock protein BMAL1, as well as physical interaction was reported (Hogenesch et al., 1998; Wu et al., 2017). Collectively, these studies point towards an intricate interplay between oxygen rhythmicity and circadian clock control. Understanding the principles underlying daily oscillations in oxygen levels are therefore of great interest. Furthermore, since oxygen is exchanged with carbon dioxide during respiration, it raises the question regarding the profile and determinants of carbon dioxide release throughout the day.

In the current study, we employed metabolic cages to measure oxygen consumption, carbon dioxide release, food intake and locomotor activity, alongside a telemetric system to record tissue (e.g. kidney) oxygenation in freely moving animals and dissected the daily control of oxygen and carbon dioxide by circadian clocks, light-dark cycles and feeding rhythms. Specifically, we asked: (i) what regulates the daily oscillations in oxygen consumption and tissue oxygenation; (ii) are there daily rhythms in carbon dioxide release and what controls them; (iii) can these processes be disintegrated and what are the potential molecular downstream effects. We found that daily rhythms in oxygen and carbon dioxide are under circadian clock control and feeding. Importantly, day-time feeding dissociated oxygen rhythms from carbon dioxide oscillations, whereby the former followed activity and the latter was shifted and aligned with food intake. Finally, we demonstrated that changes in carbon dioxide levels, at least in cultured cells, alter clock-gene expression and phase shift the clock. Thus, supporting a potential role for carbon dioxide in phase resetting of circadian clocks in peripheral organs *in vivo*.

Results

Daily rhythms in oxygen consumption, tissue oxygenation, and carbon dioxide release

We monitored daily changes in respiration (i.e. oxygen consumption and carbon dioxide release) as well as spontaneous locomotor activity and food intake of freely moving male and female mice at high temporal resolution using metabolic cages (Figure 1). As expected from nocturnal animals, locomotor activity and food consumption were elevated during the dark phase and were accompanied by increase in oxygen consumption. The rise in oxygen consumption during the dark phase was associated with an increase in carbon dioxide release, which exhibited a similar daily profile as oxygen consumption (Figure 1A-C). A cosine fit with a 24 h period assigned similar phases (Zeitgeber (ZT) ~18) for all 4 parameters, namely oxygen consumption and carbon dioxide release, locomotor activity and food intake, for both males and females (Figure 1D). Notably, females showed elevated respiration compared to males with higher total oxygen consumption and carbon dioxide release (Figure 1C and 1E). This is consistent with the fact that smaller animals (e.g. female compared to male mice) have higher energy expenditure (Schmidt-Nielsen, 1983), which might be in part due to their elevated locomotor activity (Figure 1E). Likewise, metabolic cage analyses of male rats revealed similar daily rhythms in the above indicated parameters (Figure S1A-D). Overall these findings are in line with previous measurements of indirect calorimetry in rodents (Fischer et al., 2018; Meyer et al., 2015; Tschop et al., 2011).

Continuous recordings of tissue oxygenation (i.e. kidney and liver) from freely moving rats using a telemetric oxygen monitoring system (Figure S1E) evinced overt daily rhythms in line with our metabolic cage analyses and in agreement with previous work of others and ours (Adamovich et al., 2017; Emans et al., 2017). Remarkably, the whole animal profile of oxygen consumption was relatively square shaped (Figure 1A-C and S1A-B), while kidney and liver oxygen profiles exhibited more of a sine wave form (Figure S1E); signifying the presence of homeostatic mechanisms that mitigate the sharp changes in oxygen consumption between the resting and

active phases. Of note, due to the size and weight of the oxygen sensor, telemetric oxygen measurements are currently non-feasible in mice and were therefore limited to experiments with rats.

The presence of overt rhythms in oxygen consumption, tissue oxygenation and carbon dioxide release in animals fed *ad libitum* prompted us to examine potential determinants that drive and coordinate these rhythms.

Circadian clocks control daily rhythms in oxygen consumption and carbon dioxide release

As aforementioned, circadian clocks are implicated in daily regulation of physiology and behavior in mammals. At the cellular level, clocks control daily rhythms in mitochondrial respiration (Manella and Asher, 2016; Neufeld-Cohen et al., 2016; Peek et al., 2013). We, therefore, examined the role of circadian clocks in control of daily oxygen consumption and carbon dioxide release. To this end we employed two different clock mutant mouse models, namely *Bmal1$^{-/-}$* and *Per1/2$^{-/-}$* mice. In the former, the positive (activator) limb of the core clock oscillator is compromised, and in the latter the negative (repressor) arm is impaired. Both clock mutant mouse models exhibit arrhythmic locomotor activity in constant dark, and circadian expression of core clock and clock-controlled genes is largely diminished (Adamovich et al., 2014; Bunger et al., 2000; Zheng et al., 2001).

Overall, under 12 h light-dark regimen both PER1/2 and BMAL1 deficient animals preserved the daily rhythmicity in oxygen consumption, and carbon dioxide release (Figure 2A-C and S2A-C for *Per1/2$^{-/-}$* and *Bmal1$^{-/-}$*, respectively) with similar phases (Figure 2D and S2D), yet the amplitude was reduced exclusively in the case of BMAL1 deficient mice (Figure 2M and S2M). Interestingly, the oxygen and carbon dioxide profiles of PER1/2 null mice mostly differed from their wild type counterparts during the dark phase, whereas a greater difference was observed in the case of BMAL1 deficient mice within the light phase (Figure 2C and S2C). This was evident

as well in the oxygen and carbon dioxide levels in the day vs. night comparison (Figure 2N and S2N).

As expected and in contrast to wild type mice, both PER1/2 and BMAL1 deficient mice showed no rhythmicity in locomotor activity, as well as feeding behavior, in constant dark. Notably, oxygen consumption and carbon dioxide release were relatively constant throughout the day in both clock mutant mouse models (Figure 2E-G and S2E-G for *Per1/2[-/-]* and *Bmal1^{-/-}*, respectively). In accordance, the amplitude of the 24 h cosine fit was lower than the standard deviation of the data for both PER1/2 and BMAL1 deficient mice, therefore considered as non-rhythmic, and a phase was not assigned (Figure 2H and 2M for *Per1/2[-/-]* and Figure S2H and S2M for *Bmal1^{-/-}*).

Based on the above-described experiments conducted with two different clock deficient mouse models, we concluded that oxygen and carbon dioxide rhythms are circadian clock controlled, plausibly through driving rhythmic activity and feeding behavior.

Time-restricted feeding restores daily oxygen and carbon dioxide rhythms in the absence of a functional clock

To test the specific role of cyclic feeding behavior in driving rhythms in oxygen consumption and carbon dioxide release we imposed time-restricted feeding during the subjective night on PER1/2 and BMAL1 deficient mice housed in constant dark. Overall, time-restricted feeding restored rhythmicity in oxygen consumption, and carbon dioxide release with similar phases as in wild type mice in both clock mutant mouse models (Figure 2I-L and S2I-L for *Per1/2[-/-]* and *Bmal1^{-/-}*, respectively). Here again, the amplitude was shallower in BMAL1 compared to PER1/2 deficient and wild type mice (Figure 2M and S2M).

Next, we examined the potency of light-dark versus feeding-fasting cycles as drivers of daily rhythms in oxygen consumption, and carbon dioxide release in the absence of a functional clock. Our analysis evinced that in both clock mutant mouse models, namely PER1/2 and BMAL1

deficient mice, the effects of light-dark cycles and feeding-fasting rhythms are comparable. Both Zeitgebers restored rhythmicity in oxygen consumption and carbon dioxide release with similar phases and amplitudes (Figure 2L-M and S2L-M). Yet we noted a trend toward higher amplitude in carbon dioxide rhythms upon time-restricted feeding compared to light-dark cycles irrespective of the clock mutant mouse model tested. Overall, we did not observe any significant changes in total oxygen consumption, carbon dioxide release, activity and food consumption throughout the day between the different mouse strains regardless of the light or feeding regimen tested above (Figure 2O and S2O).

Taken together, our findings indicate that daily oscillations in oxygen and carbon dioxide are under circadian clock control and that feeding rhythms are sufficient to drive their rhythmicity independently of the clock. Since in both clock mutant mouse models cyclic feeding behavior carried a more pronounced effect on carbon dioxide rhythms rather on oxygen oscillations, we posited that carbon dioxide rhythms might be dominated by feeding behavior.

Feeding time alters daily rhythms in oxygen consumption, tissue oxygenation, and carbon dioxide release

To delineate the specific role of feeding schedule on oxygen and carbon dioxide rhythms, wild type animals were fed either exclusively during the dark (night-fed) or the light phase (day-fed). Similar to animals fed *ad libitum*, night-fed animals exhibited robust daily rhythms in oxygen consumption, and carbon dioxide release (Figure 3A, 3B and 3D), with comparable phases (Figure 3E) yet higher amplitude in night-fed animals (Figure 3F). Notably, in night-fed animals the night/day ratio in oxygen consumption was slightly elevated compared to mice fed *ad libitum* (from 1.17 in *ad libitum* to 1.24 in night-fed), whereas a more pronounced effect was observed for carbon dioxide (from 1.28 in *ad libitum* to 1.48 in night-fed), (Figure 3G). Thus, further supporting the prominent effect of feeding on carbon dioxide. The total daily oxygen consumption and carbon dioxide release were similar under both feeding regimens (Figure 3G).

By contrast, day-time feeding resulted in pronounced effects on the daily profiles of oxygen, carbon dioxide, and locomotor activity in comparison to night-time or *ad libitum* feeding. First, instead of a single peak in night-time or *ad libitum* feeding, we now observed two peaks, both for oxygen consumption and locomotor activity, one in the beginning of the light phase and a second peak at the interphase between the light and dark phases (Figure 3C-3D). Consistently, a 24 h cosine fit that was used to compute the phase and amplitude was either lower than (locomotor activity) or very close (oxygen consumption) to the standard deviation of the data (Figure 3F). Therefore, in the absence of a reliable cosine fit, we refrained from plotting the computed phase in Figure 3E and thereafter (i.e. Figure S3, 4, S4, and 5). Second, the total oxygen consumption during the day-time and night-time was even (Figure 3G). Third, the daily profile of carbon dioxide largely differed from the one detected for oxygen and locomotion. We did not observe two distinct peaks for carbon dioxide release but rather a complete inversion with elevated levels during the light-phase compared to the night-phase in accord with feeding time (Figure 3C, 3D and 3G). The effect of day-time feeding on oxygen consumption and carbon dioxide release was preserved in female mice (Figure S3A-D) suggesting that the phenomenon is gender-independent. Furthermore, the additional sharp peak in oxygen consumption upon day-time feeding at the beginning of the light phase was evident as well in kidney oxygenation as measured in freely moving rats using our telemetric tissue oxygen monitoring system (Figure S3E).

Taken together, our results suggest that feeding schedule shape the daily profiles of oxygen consumption and tissue oxygenation, as well as carbon dioxide release. Remarkably, day-time feeding dissociated oxygen rhythms from carbon dioxide oscillations, whereby oxygen rhythms followed locomotor activity and carbon dioxide oscillations were shifted and aligned with food intake.

The effects of day-time feeding on oxygen and carbon dioxide rhythms are light-independent and do not require a functional molecular clock

The striking differential effect of day-time feeding on oxygen and carbon dioxide rhythms prompted us to examine potential underlying determinants, namely light-dark and the molecular clock. Since light carries a dominant effect on rodents' behavior and mice normally limit their locomotor activity to the dark and minimize it during the light phase, we first asked whether light plays a role in the observed pattern of oxygen and carbon dioxide rhythms upon day-time feeding. Wild type mice were housed either under 12 h light-dark regimen or in constant dark and fed exclusively during the light phase or the subjective light phase in the case of constant dark (Figure 4). The two peaks in oxygen consumption and activity and the inversion in carbon dioxide release were preserved in constant dark, and their profiles resembled to mice housed under light-dark regimen (Figure 4A-D).

Next, we asked whether these effects are dependent on the molecular clock. To this aim, we employed the $Per1/2^{-/-}$ and $Bmal1^{-/-}$ clock mutant mice, and fed them either exclusively during the dark or the light phase (Figure 5A-D and S4A-D for $Per1/2^{-/-}$ and $Bmal1^{-/-}$, respectively). Similar to wild type mice (Figures 3 and 4), and in contrast to night-fed animals, PER1/2 and BMAL1 deficient mice exhibited two peaks in oxygen consumption and locomotor activity, and inversion in carbon dioxide release upon day-time feeding.

We concluded that the effects of day-time feeding on oxygen and carbon dioxide rhythms are light-independent and do not require a functional molecular clock.

The effects of day-time feeding on oxygen and carbon dioxide rhythms are immediate and sustained

Hitherto, we identified a prominent effect of day-time feeding on oxygen and carbon dioxide rhythms during days 3 to 5 of day-time feeding (Figures 3-5, S3, and S4). Previous studies that addressed the effect of day-time feeding on circadian clock gene expression evinced that day-

time feeding uncouples peripheral clocks (e.g. liver, kidney) from the master clock in the SCN region of the brain and that this effect is gradual and takes about a week to be completed (Asher et al., 2010; Damiola et al., 2000; Mukherji et al., 2015; Saini et al., 2013; Stokkan et al., 2001). Hence, it raised the intriguing question whether the observed alterations in oxygen and carbon dioxide rhythms are the direct consequence of changes in feeding, that in turn modulate rhythmic gene expression in peripheral tissues or vice versa, namely feeding alters rhythmic gene expression and consequently affects daily respiration. To address this chicken and egg question, we set out to perform a kinetic experiment in which mice were continually monitored in metabolic cages for 16 consecutive days; in the first 3 days, mice were night-fed then on day 4 food was impeded and subsequently mice were shifted to day-time feeding for 9 days and finally food was provided *ad libitum* for 3 days (Figure 6). Notably, during the 24 h of food deprivation carbon dioxide release, in contrast to oxygen consumption, did not reach the same level as in night-fed animals, but rather gradually dropped (Figure S5A), consistent with the prominent effect of food intake on carbon dioxide levels.

Upon 8 days of day-time feeding we observed a complete inversion of circadian clock gene expression in the liver in agreement with previous reports (Asher et al., 2010; Damiola et al., 2000; Mukherji et al., 2015), (Figure S5B). Our analyses showed that the two peaks in oxygen consumption and spontaneous activity as well as the shift in carbon dioxide release appear already from the first day of day-time feeding (Figure 6A) and are preserved for 9 consecutive days with similar peak times (Figure 6B) and day versus night levels (Figure 6C). Likewise, once mice are shifted back from day-feeding to *ad libitum* the response is immediate. Hence, we concluded that the phenomenon is not transient but rather sustained and that it appears instantly upon day-time feeding. It is, therefore, conceivable that the observed changes in respiration upon day-time feeding are upstream to the gradual changes in circadian clock gene expression in peripheral tissues and might even play a role in this process.

Carbon dioxide levels affect core clock gene expression and phase shift the clock

The sustained and immediate effect of day-time feeding on respiration, in particular the prominent and rapid inversion of carbon dioxide rhythms in alignment with feeding time, prompted us to examine whether carbon dioxide levels affect core clock gene expression and consequently phase shift the clock.

To this end, we employed cell culture assays in which carbon dioxide levels can be well controlled. We first tested the effect of lowering carbon dioxide levels on the expression of the core clock genes (Figure 7A). Time course analysis revealed that moving from standard growth conditions of 5% carbon dioxide to 1% affect the expression of several core clock genes in cultured cells, in particular *Rev-Erbα, Per2*, *Cry1* and *Dbp*. Interestingly, the transcript levels of both *Per2* and *Cry1* were up-regulated, whereas *Rev-Erbα,* and *Dbp* were down-regulated. *Clock*, *Bmal1*, as well as *Per1* and *Cry2* were relatively unaffected. The effect of carbon dioxide on core clock gene expression encouraged us to test whether these changes are sufficient to elicit a phase shift. To this aim, we used NIH3T3 cell line that stably expresses a luciferase reporter gene under the control of the *Bmal1* promoter (*Bmal1*-luciferase reporter) (Nagoshi et al., 2004). A shift from 5% to 1% resulted in phase advance of almost 5 h (Figure 7B). Notably, a pulse of 10 h of 1% carbon dioxide had a more pronounced effect with phase shift of ~8 h (Figure 7C). This differential effect incited us to perform a Phase Response Curve (PRC). The PRC illustrates the relationship between time of administration (relative to the internal circadian clock) and the magnitude of the treatment's effect on circadian phase. Hence, we exposed the cells to a 2 h pulse of low carbon dioxide (i.e. 1%) during different time points within the circadian cycle and analyzed the phase shift based on the bioluminescence recording of the NIH3T3:*Bmal1*-luciferase cells (Figure S6). Based on the data depicted in Figure S5 we generated a PRC, in which the magnitude of the phase shift was plotted against circadian time (Figure 7D). Our PRC evinced that the magnitude and the direction of the phase shift depends on the time of the day

and qualifies as type-1 in chronobiological parlance. It is similar to the response observed for light-induced phase shifts of circadian behavior in laboratory animals. Of note, we did not observe any significant effect on the period length in none of the above described experiments (i.e. 5% to 1% shift, 10 h or 2 h pulse) between the control and the treated cells (Figures 7B, 7C and S6).

In summary, our results show that changes in carbon dioxide levels are sufficient to affect the expression of several core clock genes and phase shift the clock at least in cultured cells. It is, therefore, possible that the observed change in carbon dioxide rhythmicity upon day-time feeding plays a role in the inversion of circadian clocks in peripheral tissues in this conjuncture.

Discussion

Detailed measurements of behavioral (e.g. activity, feeding) alongside physiological parameters (e.g. respiration) are evidently necessary for uncovering regulatory principles that dictate their daily changes both endogenously and in response to environmental changes (Acosta-Rodriguez et al., 2017). In this conjuncture, there is a growing usage of metabolic cages that continuously monitor oxygen consumption and carbon dioxide release for energy production calculations (i.e. indirect calorimetry) together with activity and food intake to follow daily physiology and behavior (Mina et al., 2018; Tschop et al., 2011).

In this study, we monitored oxygen consumption, carbon dioxide release, food intake and spontaneous locomotor activity, alongside tissue oxygenation in freely moving animals to dissect the daily control of oxygen and carbon dioxide rhythms by circadian clocks, light-dark and feeding cycles. Our analyses of two different circadian clock mutant mouse models (i.e. PER1/2 and BMAL1 null mice) under free running condition (i.e. constant dark) strongly suggest that circadian clocks control daily rhythms in oxygen and carbon dioxide. It is likely that the effect is indirect and is the consequence of rhythmic activity and feeding behavior that are driven by the circadian clock. Both PER1/2 and BMAL1 null mice exhibited overt rhythms in respiration

when housed under 12 h light-dark regimen, yet BMAL1 deficient mice showed a shallower amplitude. The BMAL1 null mice were previously shown to exhibit early aging and age-related pathologies (Kondratov et al., 2006), however in our experiments with ~3 months old mice, we did not observe any overt signs of aging and their total oxygen consumption, carbon dioxide release, activity and feeding were comparable to the wild type control group. It is possible that the abrogated circadian amplitude in the aforementioned parameters precedes and even plays a role in the accelerated aging phenotype of these mice.

Intriguingly, the clock mutant CRY1/2 null mice were reported to show no rhythmicity in their Respiratory Exchange Rate (RER). (i.e. the ratio between carbon dioxide release and oxygen consumption) under light-dark regimen (Vollmers et al., 2009). It is unclear whether the observed difference between CRY1/2 and the PER1/2 and BMAL1 deficient mice in our experiments reflects a true biological effect, or rather stems from technical factors (e.g. light intensity/spectrum, statistical analysis). Of note, the statistical toolkit for circadian analyses mostly relies on fitting a cosine curve, and hence is not ideal for analysis of square shaped rhythms that are often observed for oxygen consumption, carbon dioxide release, activity and food intake. Future studies are expected to clarify whether CRY1/2 deficient mice indeed differ from BMAL1 and PER1/2 null mice in regard to their rhythmic respiration under light-dark cycles.

Our experiments evinced that both light-dark cycles and feeding-fasting rhythms are sufficient and as effective in driving rhythmicity in oxygen consumption and carbon dioxide release with comparable phases and amplitudes in the absence of a functional clock (i.e. BMAL1 and PER1/2 null mice). Similar effect of feeding-fasting cycles was observed with CRY1/2 deficient animals (Vollmers et al., 2009). Notably, mice that are on high fat diet loss their rhythmic feeding behavior and exhibit altered circadian gene expression (Eckel-Mahan et al., 2013; Hatori et al., 2012; Kohsaka et al., 2007). Upon this nutritional challenge, time-restricted feeding restores rhythms in RER and oxygen consumption as well independently of the circadian clock (Chaix

et al., 2018). It, therefore, appears that light-dark cycles can drive rhythmic feeding behavior, which in turn generates rhythms in oxygen and carbon dioxide independently of the clock. Another nutritional challenge that was recently shown to alter circadian gene expression is ketogenic diet (Tognini et al., 2017). Interestingly, ketogenic diet, similar to high fat diet, diminishes daily RER rhythmicity, however it does not affect the animals rhythmic feeding behavior. Overall, these studies highlight the importance of the diet composition on daily respiration.

Consistent with our model that carbon dioxide oscillations are directed by feeding, we noted a trend of higher amplitude in carbon dioxide oscillations upon feeding rhythms compared to light-dark cycles and a sharp decline over 24 h food deprivation. Furthermore, day-time feeding dissociated oxygen rhythms from carbon dioxide oscillations, whereby the former followed activity and the latter was shifted and aligned with food intake. The bimodal profile of oxygen and activity and the inversion in carbon dioxide were independent of light-dark cycles and the circadian clock. Thus, they appear as a direct response to food intake. It is plausible that the sharp spike in oxygen and activity at the dark-light transition upon day-time feeding most likely reflects the sharp bout of activity and food consumption right at the start of the unexpected food availability.

In the above-described experiments, both mice and rats were housed in standard animal facilities with ambient temperature of $\sim 22^0C$. This was set primarily to match the comfort requirement of the experimenters and the animal facility staff, but is far from the thermoneutral temperature of rodents, which is $\sim 30^0C$. Consequently, rodents increase their energy expenditure in order to maintain their homeothermy, and as a result the observed oxygen consumption levels are higher than expected for a typical mammal of their size (Schmidt-Nielsen, 1983). Despite the non-thermoneutrality conditions and in agreement with works of others and ours, mice show a clear diurnal pattern of energy expenditure, with night-time levels being higher than day-time (Fischer et al., 2018). Hence, although there is an ongoing debate concerning optimal housing

temperatures for mice to better model human metabolism/disease (Gaskill and Garner, 2014; Karp, 2012; Speakman and Keijer, 2012) the non-thermoneutral conditions that were maintained throughout this entire study, are widely common and most likely did not affect the presence of daily rhythms. It is noteworthy that since oxygen consumption in mammals per unit body mass decreases regularly with increasing body size (Schmidt-Nielsen, 1983) it explains the difference in the magnitude of oxygen consumption between male and female mice (females are smaller than males), as well as between mice and rats (the latter are bigger).

While the master clock in the SCN region of the brain is entrained by light-dark cycles, feeding rhythms serve as the dominant timing cue for peripheral clocks (Dibner et al., 2010). The adaptation of peripheral clocks to changes in feeding regimen (e.g. from night-fed to day-fed) is gradual and normally takes several days. The process most likely involves a wide variety of signaling pathways that work in concert. Therefore, kinetic experiments turned out to be instrumental in addressing potential mechanisms that are implicated in the processes (Asher and Schibler, 2011). The immediate and sustained response of respiration to changes in feeding regimen hinted towards potential involvement of blood gases in phase resetting of peripheral clocks. Recently we showed that oxygen can reset peripheral clocks through HIF1α (Adamovich et al., 2017). Here we examined the potential of carbon dioxide to phase shift the clocks. At least in cultured cells, carbon dioxide levels altered core clock gene expression and induced a prominent phase shift. Thus, supporting a potential role for carbon dioxide in phase resetting of circadian clocks in peripheral organs *in vivo*. Since carbon dioxide levels play a major role in acid-base balance, it is possible that the effect of carbon dioxide on the clock is mediated via changes in pH levels. Indeed, it was previously reported that pH levels can phase shift the clock (Kon et al., 2008). Interestingly, cells can sense changes in carbon dioxide levels irrespectively of changes in pH (Cummins et al., 2014), this raises the intriguing possibility that similar to carbon monoxide (Dioum et al., 2002; Klemz et al., 2017), carbon dioxide directly and independently of pH affects the clocks. Future studies are expected to shed light on the potential

molecular mechanism implicated in the effect on carbon dioxide on the core clock circuitry and their interplay with other resetting cues such as oxygen and temperature. These mechanisms are probably not mutually exclusive and might very well act together to phase shift clocks in peripheral organs.

Overall, the current study dissects the function of circadian clocks, light-dark and feeding cycles in control of oxygen and carbon dioxide rhythms and supports a potential role for carbon dioxide in phase resetting of peripheral clocks. Our findings that day-time feeding dissociate oxygen rhythms from carbon dioxide oscillations are potentially of clinical relevance. Namely, feeding-activity misalignment results in oxygen-carbon dioxide mismatch, and likely affects acid-base balance. These homeostatic imbalances potentially plays a role in pathologies associated with altered feeding/activity rhythms, for example in shift-workers (Brum et al., 2015).

Limitations of the Study

Our study mostly relies on metabolic cage measurements of oxygen consumption, carbon dioxide release, activity and feeding behavior, alongside telemetric measurements of tissue oxygenation in freely moving animals. However, the latter, as above mentioned, are limited to rats, and currently due to technical limitations cannot be performed with mice (of particular interest are the different clock mutant mouse models). Moreover, telemetric measurements of tissues' carbon dioxide levels in freely moving animals are currently technically unfeasible. We show that oxygen and carbon dioxide rhythms are circadian clock-controlled, most likely indirectly through rhythmic activity and feeding behavior that are driven by the clock. Yet, we cannot exclude the possibility that peripheral clocks in various tissues are directly implicated as well. Lastly, we provide evidence that oxygen rhythms are dominated by activity while carbon dioxide oscillations align with food intake. It is practically impossible to completely uncouple activity from feeding, likewise oxygen consumption from carbon dioxide release. Hence, at this stage we can only point towards dominance of activity on oxygen consumption and food intake on carbon dioxide release.

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Author Contributions

Conceptualization: Y.A., B.L. and G.A; Investigation: Y.A., B.L., A.N.C., M.G. Y.K., M.H.A., A.T., and M.P.K; Visualization: Y.A., B.L. and J.S.; Data Curation: J.S.; Software: J.S.; Funding: G.A.; Writing–Review & Editing: Y.A., B.L. and G.A.

Declaration of Interests

The authors declare no competing interests.

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Main Figure Titles and legends

Figure 1. Daily rhythms in oxygen consumption and carbon dioxide release. Wild type male (M) and female (F) mice were housed under 12 h Light-Dark cycles. Metabolic cages were used to monitor: oxygen consumption rate (VO_2) (blue); carbon dioxide release rate (VCO_2) (red); spontaneous locomotor activity (green) and food intake (black). Data are presented as the mean ± SEM of 3 mice, recorded at 15 min intervals for 3 consecutive days (**A** and **B** males and females, respectively), alongside the mean ± SEM of three days' average of each individual mouse (**C**), the calculated phase and amplitude (Amp) based on 24 h cosine fit analysis (**D**), and the total (mean ± SEM) for the Light phase, Dark phase and the entire Day (**E**). Zeitgeber Time (ZT); Gray shadings represent the dark phases. t-test, *$p < 0.05$, **$p<0.01$, ***$p<0.001$, n.s (non-specific). See also Figure S1.

Figure 2. Time-restricted feeding restores daily oxygen and carbon dioxide rhythms in *Per1/2$^{-/-}$* mice in Dark-Dark. Wild type (WT) and *Per1/2$^{-/-}$* (KO) mice were housed under 12 h Light-Dark (**A-D**), or Dark-Dark (**E-L**), and fed either *ad libitum* (*Ad lib*) (**A-H**), or exclusively during the subjective night (tRF) (**I-L**). The same cohort of mice was used for all conditions. Metabolic cages were used to monitor: oxygen consumption rate (VO_2) (blue); carbon dioxide release rate (VCO_2) (red); spontaneous locomotor activity (green) and food consumption (black). Data are presented as the mean ± SEM of 3 mice, recorded at 15 min intervals for 3 consecutive days (**A**, **B**, **E**, **F**, **I**, and **J**), alongside the mean ± SEM of three days' average of each individual mouse (**C**, **G**, and **K**), the calculated phase (**D**, **H**, and **L**) and amplitude (mean ± SEM) based on 24 h cosine fit analysis (**M**), the total (mean ± SEM) for the Light and Dark phases (**N**), and the total per day (mean ± SEM) (**O**). Zeitgeber Time (ZT), Circadian time (CT); For **A-D**; Gray shadings represent the dark phases. For **E-L**; Dark and light gray shadings represent the subjective dark and light phases, respectively. L/D: 12 h light-dark; D/D: constant dark. Orange bars mark times of food availability. t-test, *$p < 0.05$, **$p<0.01$, n.s (non-specific). # marks cases

in which amplitude divided by the S.D. of the data was below 1, hence considered not to exhibit 24 h rhythmicity.

Figure 3. The effect of feeding-time on daily oxygen and carbon dioxide rhythms in wild type mice. Wild type mice were housed under 12 h light-dark regimen and either fed *ad libitum* (*Ad-lib*) (**A**), or during the dark phase (Days 2-4 upon Night-feeding), (Night-Fed) (**B**), or the light phase (Days 3-5 upon day-feeding), (Day-Fed) (**C**). The same cohort of mice was used for all 3 conditions. Metabolic cages were used to monitor: oxygen consumption rate (VO_2) (blue); carbon dioxide release rate (VCO_2) (red); spontaneous locomotor activity (green) and food consumption (black). Data are presented as the mean of 7 mice, recorded at 15 min intervals for 3 consecutive days, (**A**, **B**, and **C**), alongside the mean ± SEM of three days' average of each individual mouse (**D**), the calculated phase (**E**), the amplitude (mean ± SEM) based on 24 h cosine fit analysis (**F**), and the total (mean ± SEM) for the Light phase, Dark phase and the entire day (**G**). Zeitgeber Time (ZT); Gray shadings represent the dark phases. Orange bars mark times of food availability. t-test, **$p<0.01$, ***$p<0.001$, n.s (non-specific). # marks cases in which amplitude divided by the S.D. of the data was below 1, hence considered not to exhibit 24 h rhythmicity. See also Figure S3.

Figure 4. The effect of feeding-time on daily oxygen and carbon dioxide rhythms is light-dark cycles independent. Wild type mice were housed under 12 h Light-Dark (L/D) (**A**), or Dark-Dark (D/D) (**B**), and fed exclusively during the light phase (subjective light phase in Dark-Dark). The same cohort of mice was used for both conditions. Metabolic cages were used to monitor: oxygen consumption rate (VO_2) (blue); carbon dioxide release rate (VCO_2) (red); spontaneous locomotor activity (green) and food consumption (black). Data are presented as the mean of 5 mice, recorded at 15 min intervals for 3 consecutive days, (**A**, and **B**), alongside the mean ± SEM of three days' average of each individual mouse for the L/D condition, and the

mean ± SEM of two days' average of each individual mouse for the D/D condition (due to a technical problem marked with [*]) (**C**), and the total (mean ± SEM) for the Light phase, Dark phase and the entire day (**D**). Zeitgeber Time (ZT); Circadian Time (CT); For **A**; Gray shadings represent the dark phases. For **B**; Dark and light gray shadings represent the subjective dark and light phases, respectively. Orange bars mark times of food availability. t-test, **p<0.01, n.s (non-specific). [*] marks a transient technical problem in the recordings (automated feeders were open for 2 h at the wrong time).

Figure 5. The effect of feeding-time on daily oxygen and carbon dioxide rhythms in *Per1/2⁻/⁻* mice. *Per1/2⁻/⁻* mice were housed under 12 h light-dark regimen and fed exclusively during the dark phase (Days 2-4 upon Night-feeding), (Night-Fed) (**A**), or the light phase (Days 3-5 upon day-feeding), (Day-Fed) (**B**). The same cohort of mice was used for both conditions. Metabolic cages were used to monitor: oxygen consumption rate (VO_2) (blue); carbon dioxide release rate (VCO_2) (red); spontaneous locomotor activity (green) and food consumption (black). Data are presented as the mean of 3 mice, recorded at 15 min intervals for 3 consecutive days, (**A**, and **B**), alongside the mean ± SEM of three days' average of each individual mouse (**C**), and the total (mean ± SEM) for the Light phase, Dark phase and the entire day (**D**). Zeitgeber Time (ZT); Gray shadings represent the dark phases. Orange bars mark times of food availability. t-test, *p < 0.05, ***p<0.001, n.s (non-specific). See also Figure S4.

Figure 6. The effects of day-time feeding on oxygen and carbon dioxide rhythms are immediate and sustained. Wild type mice were continually monitored in metabolic cages for 16 consecutive days; in the first 3 days mice were night fed, then on day 4 food was impeded and subsequently mice were shifted to day-time feeding for 9 days, next food was provided *ad libitum* for 3 days. Metabolic cages were used to monitor: oxygen consumption rate (VO_2) (blue); carbon dioxide release rate (VCO_2) (red); spontaneous locomotor activity (green) and food

consumption (black). Data are presented as the mean of 3 mice, recorded at 15 min intervals for 3 consecutive days (**A**), alongside the peak detection in each day (the last peak in VCO_2 data was not detected due to edge effect) (**B**), and the total (mean ± SEM) for the Light phase and Dark phase for each day (**C**). Zeitgeber Time (ZT); Gray shadings represent the dark phase. The circle size represents the prominence of the peak. The arrow indicates the first day of day-time feeding. Orange bars mark times of food availability. t-test, *p < 0.05, ** p<0.01 ***p<0.001. [*] marks an unexpected technical problem in the activity recordings for 9 h (another set of recordings for the 24 h of food deprivation is presented in Figure S5A). See also Figure S5.

Figure 7. Carbon dioxide levels affect core clock gene expression and phase shift the clock. (**A**) NIH3T3 cells were maintained at 5% CO_2. After 3 days in culture cells were either shifted to 1% CO_2 or kept at 5% CO_2 as control. Cell were harvested at indicted times; total RNA was prepared and transcript levels of clock genes were determined by quantitative real-time PCR and presented as fold change relative to the lowest value. (**B**) Bioluminescence recordings of NIH3T3 stably expressing *Bmal1*-luciferase reporter that were cultured either at 5% CO_2 or shifted to constant 1% CO_2, alongside bar graph representation of the average circadian period ± SEM. (**C**) Bioluminescence recordings of NIH3T3 stably expressing *Bmal1*-luciferase reporter that were cultured either at 5% CO_2 or were exposed to a 10 h pulse of 1% CO_2, alongside bar graph representation of the average circadian period ± SEM. (**D**) Phase Response Curve in response to 2 h pulses of 1% CO_2. The x axis indicates the circadian time at the beginning of the pulse. The y axis indicates the phase shift resulted from the pulse. The bioluminescence recordings for each pulse are presented in Figure S5. Circadian Time (CT), n.s (non-specific). See also Figure S6.

STAR METHODS

CONTACT FOR REAGENT AND RESSOURCE SHARING

Further information and requests for resources and reagents should be directed to and will be fulfilled by the Lead Contact, Gad Asher (gad.asher@weizmann.ac.il)

EXPERIMENTAL MODEL DETAILS

Animals

All animal experiments and procedures were conducted in conformity with the Weizmann Institute Institutional Animal Care and Use Committee guidelines, the Ben-Gurion University of the Negev Animal Use and Care Committee protocol number: IL-77-11-2015 and the UK Home Office (Scientific Procedures) Act (1986) with project approval from the ethics of research committee, University of Bristol. For metabolic cages analyses, we used three months old Wistar rats and C57BL/6 wild type, *Per1/2$^{-/-}$* (Zheng et al., 2001) back crossed to C57BL/6 and C57BL/6 *Bmal1$^{-/-}$* (Bunger et al., 2000) mice. For telemetric oxygen monitoring three months old Wistar rats were used. Animals were housed under 12 h light-dark regimen or constant dark and fed either *ad libitum* or exclusively during the dark or light phase as indicated. The temperature in the housing cages and metabolic cages was maintained at 22°C. ZT0 corresponds to the time lights were turned ON and ZT12 to the time lights were turned OFF in the animal facility.

METHOD DETAILS

Metabolic cages

Mice and rats' oxygen consumption rate, carbon dioxide release, spontaneous locomotor activity and food consumption were simultaneously monitored for individually housed mice using the Phenomaster metabolic cages (TSE System) and Promethion metabolic cages (Sable Systems International), respectively. Before each experiment animals were trained for several days in the metabolic cages to enable them to properly adjust to the new housing conditions. A period of 3-7 days acclimatization was taken before the actual recordings. Data was collected at 15 min

intervals for TSE, and 1 min for Sable system. All animals presented per figure were recorded together side by side to allow adequate comparisons. The light schedule in the metabolic cages was maintained as in the animals' home cages. Fluorescent light with intensity of 100 lux was applied for the metabolic cage recordings during the light phase.

Time-restricted feeding

The same cohort of animals was fed according to the different feeding schedule per figure as indicated in the figure legend. Transition from *ad libitum* to night-time feeding was done with a 12 h fasting period during the light phase, and from night to day-time feeding with a 24 h fasting period. Time-restricted feeding was performed either manually (rats), or using automated feeders (mice).

Tissue oxygenation measurement

Oxygen levels in rat kidney/liver were measured using a telemetry system as described previously (Koeners et al., 2013; Koeners et al., 2016). Telemeters (TR57Y) equipped with a carbon paste electrode (CPE, 0.27 mm in diameter) were sterilized in a 2% w/v glutaraldehyde solution for at least 4 hours or 0.55% w/v ortho-phthalaldehyde (Cidex OPA) for 30 minutes and rinsed thoroughly with sterile 0.9% w/v NaCl solution before implantation. Rats were anaesthetized with 5% v/v isofluorane in an induction box and maintained at 2–2.5 % v/v isoflurane on a heated operating table. Rats were pre-medicated with buprenorphine (30 µg/kg s.c., Temgesic, Reckitt Benckiser, Slough, UK). Under sterile conditions, the kidney/liver and aorta were exposed by laparotomy. The cables connecting the electrodes and telemeter were secured by suturing them on the adventitia of the abdominal aorta or dorsal muscles adjacent to the spine near the kidney/liver. After pre-puncturing the tissue with a 30-gauge needle, the reference electrode and CPE, were inserted and secured in place with tissue glue (Histoacryl, 1050044 B. Braun, Tuttlingen, Germany) approximately 1 mm apart from each other, while the auxiliary electrode was affixed onto the tissue surface. With the telemeter placed and secured in

the abdomen, the abdomen was closed with sutures and the rat was placed on a heated pad for at least 12 hours to recover. Post-operative analgesia was administered (buprenorphine, 3 µg/100 g rat every 8–14 h for up to 3 days or as required). All rats were allowed to recover for 1 week before experiments started. After a recovery, the rat's cage was placed on a receiver-charging unit (SmartPad TR181, Millar Inc, Tx), which received the data from, and recharged the battery of, the telemeter. Renal/hepatic O_2 levels were continuously measured in Wistar rats for several consecutive days. After the experimental period, rats were sacrificed by intraperitoneal injection of an overdose of sodium pentobarbitone (>200 mg ml-1, Euthatal, Merial, Pirbright, UK) and post-mortem O_2 values were determined for off-set correction of individual O_2 recordings

Cell lines

A stable NIH 3T3 cell line expressing *Bmal1*-luciferase (Nagoshi et al. 2004) was cultured in DMEM containing 10% FBS, 100IU/ml penicillin and 100mg/ml streptomycin at 37^0C in a humidified incubator with 5% CO_2 unless indicated otherwise.

Real-Time Bioluminescence Monitoring

Cells (0.2×10^6 cells/plate) were seeded in 35mm culture dishes and cultured for 2 days. Cell culture media was replaced to DMEM without phenol red supplemented with luciferine and cells were placed in the LumiCycle for continuous bioluminescence recordings for several consecutive days. CO_2 levels were either kept at 5% (control) or reduced to 1% for periods indicated in the main text and figure legends. Data was analyzed with the LumiCycle analysis software (Actimetrics).

Gene Expression analysis

Total RNA was extracted from mouse liver tissue using TRI reagent (Sigma Aldrich). RNA quality and quantity were measured using Nanodrop (Thermo Scientific). cDNA synthesis was performed using qScript (Quantabio). RNA expression analysis was performed by real-time PCR using SYBR green or Taqman probes with LightCycler II machine (Roche). Expression levels

were calculated using the $\Delta\Delta$cT method and normalized to the geometrical mean of three housekeeping genes: *Hprt*, *Rplp0, Tbp*. Primers and probes (Sigma) are listed in Table S1.

QUANTIFICATION AND STATISTICAL ANALYSIS

Cosine Fits

Rhythmicity analysis was done as previously described (Sobel et al., 2017) using harmonic regression. Briefly, we used the function $x(t)=A_0+A_{24}\cos(2\pi/(24\text{ h}) * (t-t_{max}))$, where A_0 is the mean signal, A_{24} the amplitude of the 24h oscillations, and t_{max} the phase (or peak time). This method is also known as the single Cosinor model (Cornelissen, 2014). Thus, using this model, we performed least squared fitting of activity, feeding, VO_2 and VCO_2 data in each experimental condition. The following exclusion criteria was used: the cosine fit, although having a significant p-value, was considered as biologically non-relevant in case the amplitude divided by the S.D. of the data was below 1 (marked with # in the relevant figures).

Peak detection

Peak detection was performed using the built-in function findpeaks() in MATLAB, on smoothed data of 1.75 h running-average window. The function parameter values that were used: minimal peak prominence of 2 standard deviations of the signal, minimal peak distance of 6 h, and minimal peak height defined as the signal mean level.

Statistical analysis

Details regarding sample size and statistical tests are indicted in the figure legends. A two-tail Student's t tests unpaired (for amplitude and total per day) or paired (for Light phase vs Dark phase) were used with $*$ for $p < 0.05$, $**$ for $p < 0.01$ and $***$ for $p < 0.001$. Data visualizations are depicted with the average of biological replicates $\pm$ standard error of the mean. Statistical significance was concluded at $p < 0.05$. Telemetric data are represented with a 3 h smoothing

window overlapped with the raw data (every minute). Statistics were calculated with R and

Excel statistical features.