Supplemental Material

Continuous light does not affect atherosclerosis in APOE*3-Leiden.CETP mice

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Supplementary Tables and Figures

Gene	Primer see	quence	Product length (bp)
36b4	Forward Reverse	5'- GGACCCGAGAAGACCTCCTT -3' 5'- GCACATCACTCAGAATTTCAATGG -3'	85
Bmal1	Forward Reverse	5'- ATGCCAAGACTGGACTTCCG -3' 5'- TGCAGAAGCTTTTTCGATCTGC -3'	180
Clock	Forward Reverse	5'- AGTTAGGGCTGAAAGACGGC -3' 5'- GGTGTGGAGGAAGGGTCTGA -3'	216
Cryl	Forward Reverse	5'- AGAGGGCTAGGTCTTCTCGC -3' 5'- GTGAGTCTGCTGACTGTCCC -3'	222
Cry2	Forward Reverse	5'- CCAGAGCACTATCCAGTGGC -3' 5'- GCGATGGCTCTCCAGTCTC -3'	154
Per1	Forward Reverse	5'- ACGGCCAGGTGTCGTGATTA -3' 5'- CCCTTCTAGGGGACCACTCA -3'	162
Per2	Forward Reverse	5'- TGTGTGCTTACACGGGTGTCCTA -3' 5'- ACGTTTGGTTTGCGCATGAA -3'	142
Reverba	Forward Reverse	5'- GTGCTTGTCTCTGCAGACCG -3' 5'- TTGGTGAAGCGGGAAGTCTC -3'	131

Table S1. Primer Sequences for qRT-PCR

Parameter	T-value	Degrees of freedom (df)	P-value	
Food intake	2.442	34	0.0200	
Fat mass	1.052	33	0.3006	
Lean mass	1.342	33	0.1887	
Organ weight - Liver	0.662	34	0.5126	
Organ weight – Spleen	1.718	34	0.0952	
Organ weight – iBAT	1.649	34	0.1083	
Organ weight – gWAT	1.190	34	0.2422	
Organ weight – sWAT	0.747	34	0.4600	
Cholesterol exposure	2.013	34	0.0521	
Bone marrow WBCs	2.765	33	0.0092	
Bone marrow RBCs	2.821	34	0.0079	
Bone marrow PLTs	1.742	34	0.0905	
Blood WBCs	4.007	34	0.0003	
Blood RBCs	0.659	34	0.5142	
Blood PLTs	2.610	34	0.0134	
Blood neutrophils	1.507	28	0.1430	
Blood lymphocytes	0.894	29	0.3787	
Blood monocytes	0.456	29	0.6518	
Blood eosinophils	0.108	29	0.9145	
Blood basophils	2.204	34	0.0344	
Atherosclerotic lesion area	0.620	33	0.5396	
Lesion severity - mild	1.552	32	0.1302	
Lesion severity - severe	1.550	32	0.1309	
Rhythm strength	2.455	24	0.0217	
Rhythm period	10.26	24	<0.0001	
Clock expression in iBAT	0.381	16	0.7082	
Bmall expression in iBAT	2.133	16	0.0487	
Reverba expression in iBAT	4.514	16	0.0004	
Cry1 expression in iBAT	3.541	16	0.0027	
Cry2 expression in iBAT	2.330	16	0.0332	
Per1 expression in iBAT	0.709	16	0.4883	
Per2 expression in iBAT	3.005	16	0.0084	
Clock expression in gWAT	2.241	16	0.0396	
Bmall expression in gWAT	0.017	16	0.9870	
Reverba expression in gWAT	3.606	16	0.0024	

Table S2. Values of statistical testing by two-tailed unpaired T test

Cry1 expression in gWAT	1.080	16	0.2961
Cry2 expression in gWAT	1.140	16	0.2711
Per1 expression in gWAT	0.645	16	0.5278
Per2 expression in gWAT	1.921	16	0.0728
Clock expression in liver	0.856	16	0.4044
Bmal1 expression in liver	0.721	16	0.4815
Reverba expression in liver	2.825	16	0.0122
Cry1 expression in liver	1.532	16	0.1452
Cry2 expression in liver	1.391	16	0.1831
Per1 expression in liver	1.087	16	0.2932
Per2 expression in liver	0.514	16	0.6146
Clock expression in aorta	0.846	15	0.4109
Bmal1 expression in aorta	1.021	15	0.3236
Reverba expression in aorta	3.198	15	0.0060
Cry1 expression in aorta	1.821	15	0.0886
Cry2 expression in aorta	0.248	15	0.8076
Per1 expression in aorta	0.983	15	0.3410
Per2 expression in aorta	2.364	15	0.0320

P-values are marked bold when statistically significant (P < 0.05).

Parameter	Statistical test including group, time and group*time interaction effects	Adjusted P- Sidak's post	
Cumulative food intake	<u>Two-way ANOVA</u> Group F (1, 6) = 1.563; P=0.258 Time F (1.260, 7.559) = 1462; P<0.0001 Interaction F (10, 60) = 2.253; P=0.026	T≈1 week T≈3 weeks T≈4 weeks T≈5 weeks T≈6 weeks T≈7 weeks T≈8 weeks T≈9 weeks T≈10 weeks T≈11 weeks T≈12 weeks	>0.999 >0.999 0.996 0.979 0.962 0.901 0.887 0.924 0.931 0.975 0.984

Body weight Triglycerides	<u>Two-way ANOVA</u> Group F (1, 34) = 0.007; P=0.934 Time F (2.752, 93.55) = 137.2; P<0.0001 Interaction F (5, 170) = 3.502; P=0.005	T=0 weeks T=2 weeks T=4 weeks T=8 weeks T=12 weeks T=14 weeks T=0 weeks	0.782 >0.999 >0.999 0.989 0.865 0.996
	Group F (1, 34) = 2.028; P=0.164 Time F (5, 162) = 16.49; P<0.0001 Interaction F (5, 162) = 2.101; P=0.068	T=2 weeks T=4 weeks T=8 weeks T=12 weeks T=14 weeks	0.848 0.057 0.909 0.985 0.501
Total cholesterol	<u>Mixed-effects models</u> Group F (1, 34) = 4.042; P=0.052 Time F (3.160, 99.84) = 44.11; P<0.0001 Interaction F (5, 158) = 2.801; P=0.019	T=0 weeks $T=2 weeks$ $T=4 weeks$ $T=12 weeks$ $T=14 weeks$	0.999 0.583 0.234 0.379 0.205 0.954
HDL cholesterol	Mixed-effects models Group F (1,33) = 0.1119; P=0.740 Time F (2, 58) = 8.369; P=0.0006 Interaction F (2, 58) = 0.967; P=0.386	T=8 weeks T=12 weeks T=14 weeks	0.930 0.807 0.765
Non-HDL cholesterol	<u>Mixed-effects models</u> Group F (1,33) = 5.771; P=0.022 Time F (1.936, 49.36) = 20.52; P<0.0001 Interaction F (2, 51) = 1.506; P=0.2315	T=8 weeks T=12 weeks T=14 weeks	0.219 0.038 0.542
Rhythm strength per day (LD vs. LD-DL)	<u>Two-way ANOVA</u> Group F (1, 11) = 1.238; P=0.290 Time F (1.409, 15.50) = 14.17; P=0.0008 Interaction F (9, 99) = 11.89; P<0.0001	T=95 days T=96 days T=97 days T=98 days T=99 days T=100 days T=101 days T=102 days T=103 days T=104 days	>0.999 >0.999 >0.999 0.172 0.003 0.179 0.938 0.974 >0.999 >0.999

P-values are marked bold when statistically significant (P<0.05).

Table S4. Values of Pearson correlation analyses

Correlation	Group	R-squared	95% Confidence interval	P-value
Atherosclerotic lesion area with rhythm strength	LD&LL LD LL	0.0372 0.3956 0.3565	-0.2189 to 0.5464 -0.9241 to 0.1360 -0.1499 to 0.7147	0.3555 0.0948 0.1602
Severe atherosclerotic lesions with rhythm strength	LD&LL LD LL	0.00044 0.02863 0.00012	-0.3856 to 0.4209 -0.6079 to 0.7808 -0.5039 to 0.4874	0.9221 0.6887 0.9679

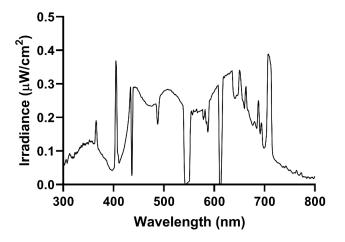


Figure S1. Spectral power distribution of the light source. Mice were housed in light-tight cabinets fitted with diffuse white fluorescent light of ~100 lux. The spectral power distribution of the light source was measured with an AvaSpec 2048-SPU (Avantes BV, Apeldoorn, The Netherlands) light meter. Adapted from Schilperoort *et al*, J Pineal Res, 2020 (DOI:10.1111/jpi.12614).

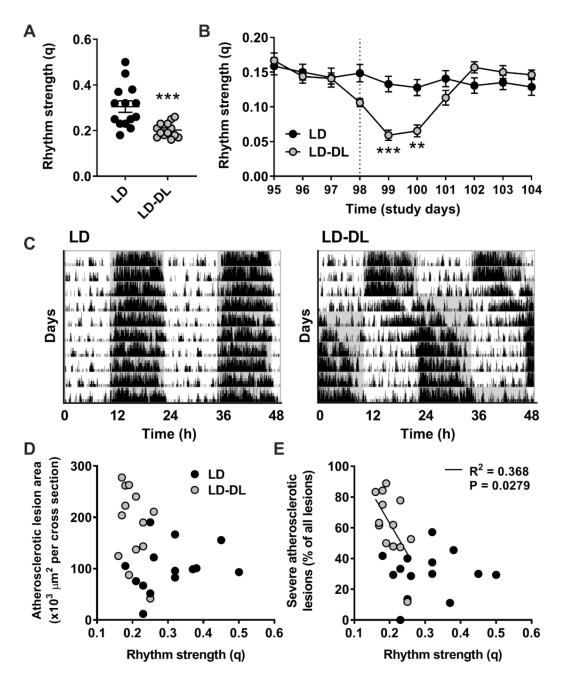


Figure S2. Shifts in light-dark cycle promote a strong acute reduction in rhythm strength, which correlates to atherosclerotic lesion severity. APOE*3-Leiden.CETP mice fed a Western-type diet were exposed to either regular light-dark cycles (LD) or weekly alternating light-dark cycles (12 h shifts; LD-DL) (n = 15/group) for 15 weeks. During week 14 and 15 of the light intervention, mice were housed individually in cages fitted with passive infrared detectors to assess behavioral activity patterns. F-periodogram analysis was performed to calculate the rhythm strength per week (A) and per day (B), and the dotted line in figure panel B indicates the day on which the LD cycle is shifted for the LD-DL group. Representative double-plotted actograms of LD and LL mice are shown (C). Rhythm strength was correlated to the atherosclerotic lesion area (D) and the relative amount of severe atherosclerotic lesions (E), and the R-squared and P-values are shown for the significant correlation between rhythm strength and severe atherosclerotic lesions within the LD-DL group, as evaluated by Pearson correlation analysis. Data are expressed as individual values or as means \pm SEM. Significance was tested by the two-tailed unpaired Student T test (A) or mixed models (B). **P<0.01, ***P<0.001 compared to the LD control group.

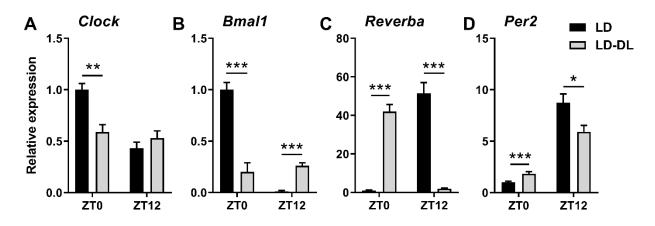


Figure S3. Expression of clock genes in the aorta is disrupted in mice exposed to shifts in lightdark cycle. APOE*3-Leiden.CETP mice fed a Western-type diet were exposed to regular light-dark cycles (LD) or weekly alternating light-dark cycles (LD-DL) (n = 18/group) for 10 weeks, after which aortas were isolated at either ZT0 or ZT12 (n = 7-8 per timepoint/group) for gene expression analysis of *Clock* (A), *Bmal1* (B), *Reverba* (C), and *Per2* (D). Data are expressed as means ± SEM. Significance was tested by two-way ANOVA. *P<0.05, **P<0.01, ***P<0.001 compared to the LD control group.

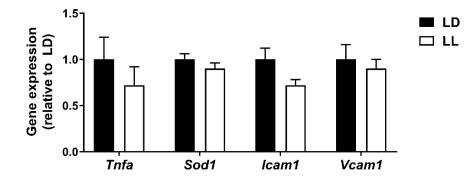


Figure S4. Expression of markers of inflammation, oxidative stress and leukocyte recruitment in the aorta is unaffected by exposure to constant light. APOE*3-Leiden.CETP mice fed a Western-type diet were exposed to LD or LL (n = 18/group) for 14 weeks, after which aortas were isolated (n = 9/group) at ZT2 for gene expression analysis of markers of inflammation (*Tnfa*), oxidative stress (*Sod1*), and leukocyte recruitment (*Icam1* and *Vcam1*). Data are expressed as means ± SEM. Significance was tested by the two-tailed unpaired Student T test.