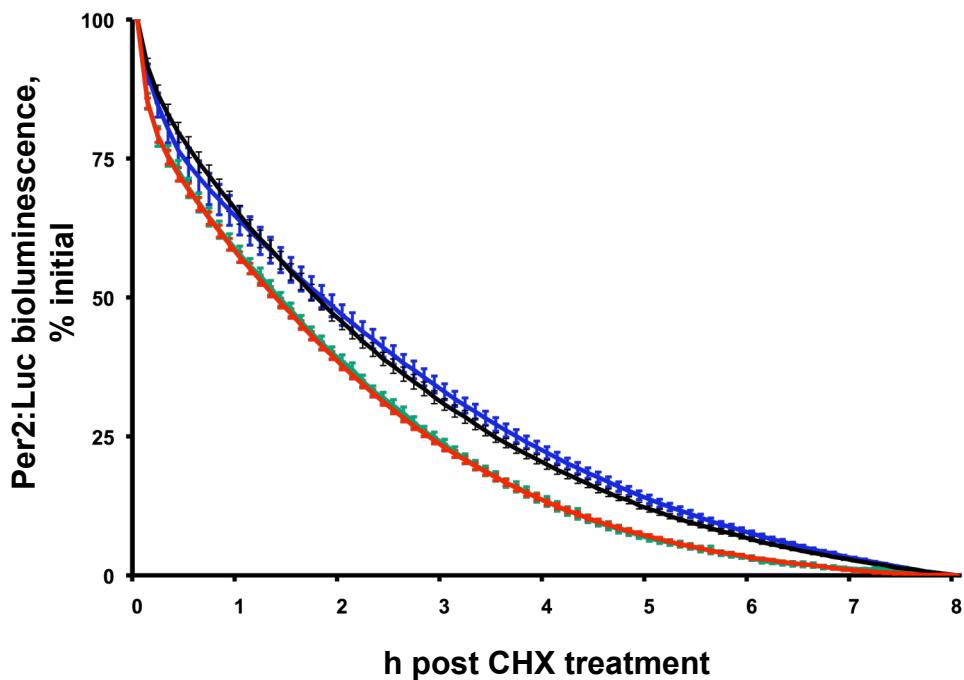


Maywood et al. Suppl. Figure S1.

Decline of Per2-mediated bioluminescence in SCN slices treated with cycloheximide: effect of $CK1\epsilon^{Tau/Tau}$ is independent of $Fbxl3^{Afh/Afh}$.

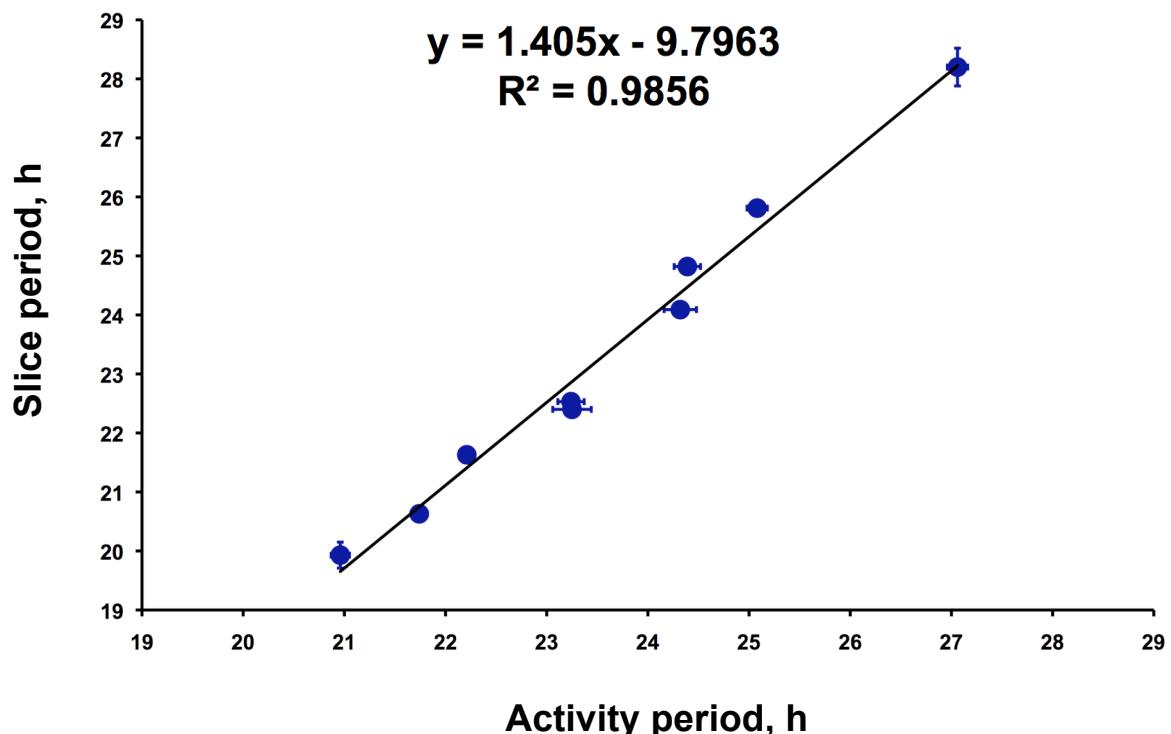
Per2:Luc bioluminescence recordings from SCN slices treated with cycloheximide (CHX) at the peak of the bioluminescence cycle. Curves were normalised to 100% for initial reading. Note comparable accelerated decline, reflecting decreased Per2 stability, in $CK1\epsilon^{Tau/Tau}$ slices with both $Fbxl3^{+/+}$ and $Fbxl3^{Afh/Afh}$ backgrounds.

Black = WT, Blue = Afh, Red = Tau, Green = Tau/Afh, (mean \pm SEM, n= 6-8 per genotype).



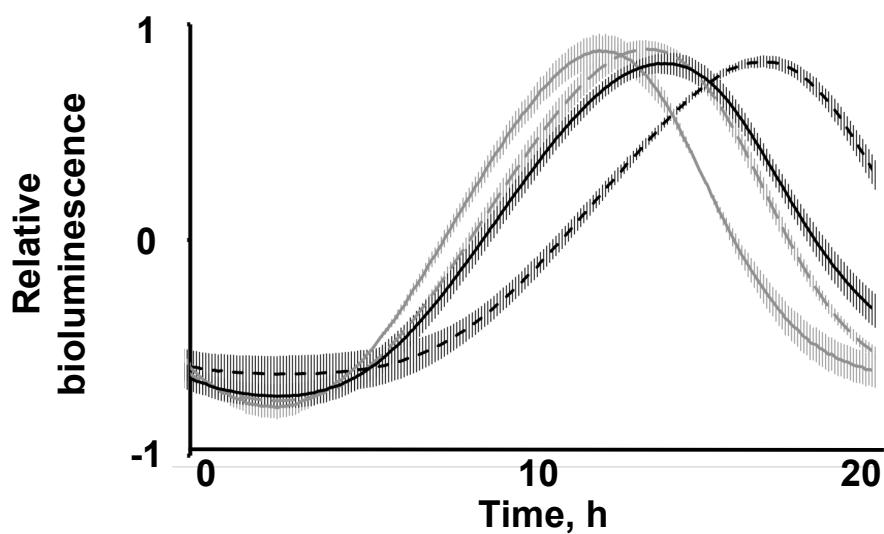
Maywood et al. Suppl. Figure S2.

Correlation between mean circadian periods of wheel-running activity and SCN bioluminescence rhythms in mice carrying single and combined *CK1ε^{Tau}* and *Fbxl3^{Afh}* mutations.



Maywood et al. Suppl. Figure S3.

Normalised plots of bioluminescence curves from wildtype (solid black), *Afh* homozygous (dashed black), *Tau* homozygous (solid grey) and double homozygous (dashed grey) SCN, aligned by nadir of bioluminescence. Curves represent mean (\pm SEM) of 5 slices per genotype.



Supplementary Table S1. Parameters of entrained and free-running circadian rhythm of wheel-running activity in wildtype and single and double mutant mice (Mean ±SEM, (n))

Genotype	Phase angle on LD, h	Period in DD, h	Periodogram Amplitude in DD
Fbxl3 ^{+/+} ::CSK1e ^{+/+}	-0.17±.07 (11)	24.32±.07 (11)	1012±140 (11)
Fbxl3 ^{+/+} ::CSK1e ^{Tau/+}	0.33±0.35 (6)	22.21±0.09 (6)	1034±212 (6)
Fbxl3 ^{+/+} ::CSK1e ^{Tau/Tau}	nd	20.96±0.22 (9)	798±180 (9)
Fbxl3 ^{Afh/+} ::CSK1e ^{+/+}	-0.52±0.10 (6)	25.08±0.13 (6)	1282±144 (6)
Fbxl3 ^{Afh/+} ::CSK1e ^{Tau/+}	-0.18±0.05 (7)	23.24±0.07 (7)	1247±133 (7)
Fbxl3 ^{Afh/+} ::CSK1e ^{Tau/Tau}	0.49±0.37 (7)	21.74±0.12 (7)	1359±123 (7)
Fbxl3 ^{Afh/Afh} ::CSK1e ^{+/+}	-0.34±0.10 (7)	27.06±0.32 (7)	989±186 (7)
Fbxl3 ^{Afh/Afh} ::CSK1e ^{Tau/+}	-0.22±0.08 (6)	24.39±0.11 (6)	1073±114 (6)
Fbxl3 ^{Afh/Afh} ::CSK1e ^{Tau/Tau}	-0.14±0.03 (10)	23.25±0.10 (10)	1047±151 (10)

Supplementary Table S2. Period, amplitude and relative amplitude error for regional bioluminescence rhythms in wildtype and double homozygous *Afh::Tau* mutant SCN slices (mean \pm SEM, n = 3 per genotype). DM lip = dorsomedial lip, the leading edge of the bioluminescent wave; DL core = dorsolateral core; core, lateral and ventral as stated.

Region	Period, h		Amplitude, cps		Relative amplitude error	
	Wild-type	Double homozygote	Wild-type	Double homozygote	Wild-type	Double homozygote
DM lip	24.19 \pm 0.28	22.71 \pm 0.14	714 \pm 197	570 \pm 242	0.083 \pm 0.019	0.120 \pm 0.023
DL core	24.32 \pm 0.34	22.93 \pm 0.15	1295 \pm 421	1247 \pm 437	0.081 \pm 0.006	0.109 \pm 0.012
Core	24.33 \pm 0.36	22.89 \pm 0.15	1090 \pm 272	1026 \pm 300	0.086 \pm 0.009	0.091 \pm 0.006
Lateral	24.35 \pm 0.38	22.91 \pm 0.17	708 \pm 266	598 \pm 205	0.082 \pm 0.007	0.090 \pm 0.009
Ventral	24.17 \pm 0.31	22.79 \pm 0.12	511 \pm 182	379 \pm 165	0.080 \pm 0.022	0.076 \pm 0.008