

Supplementary Online Material

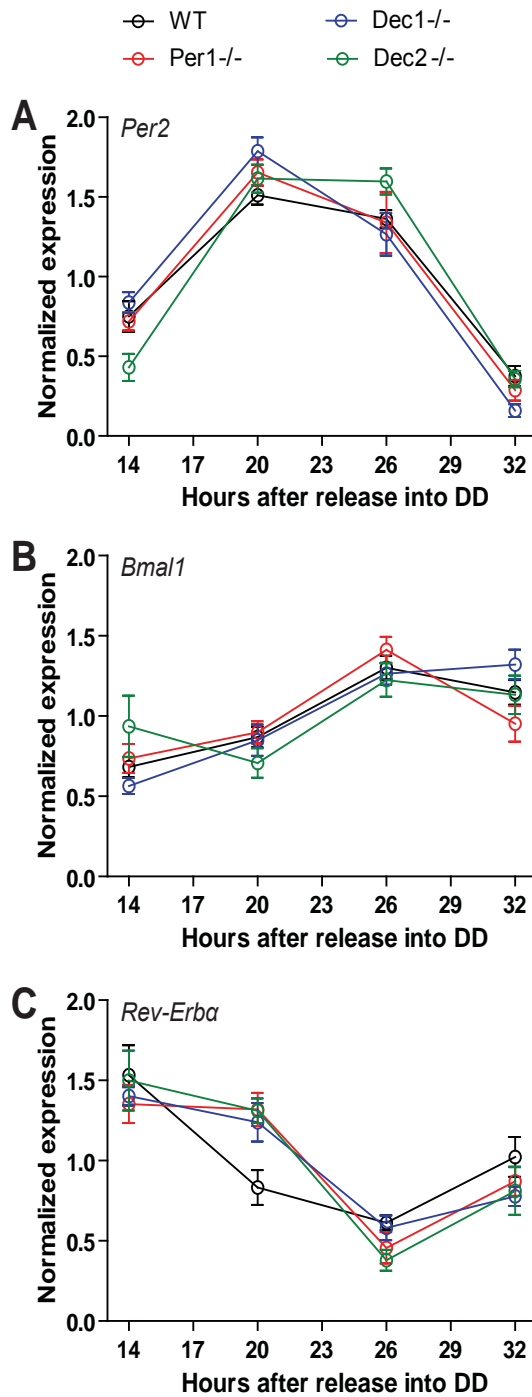
Genetic interaction of *Per1* and *Dec1/2* in the regulation of circadian locomotor activity

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Bode et al. Figure S1. In situ hybridization (ISH) profiles of clock gene expression in the SCN. Normalized expression of (A) *Per2*, (B) *Bmal1* and (C) *Rev-Erba* mRNA in the SCN in DD of wild-type (WT; black line), *Per1*^{-/-} (red line), *Dec1*^{-/-} (blue line) and *Dec2*^{-/-} (green line) mice. Data are normalized to the average expression in wild-types and double plotted for visual clarity. All values are means \pm SEM (n = 3).

Bode et al. Table S1: Statistics of behavioral paradigms

Tested parameter	Genotype comparison	↓ → Per1 ^{-/-}	Dec1 ^{-/-}	Dec2 ^{-/-}	Per1 ^{-/-} Dec1 ^{-/-}	Per1 ^{-/-} Dec2 ^{-/-}	Dec1 ^{2/-}	Per1 ^{-/-} Dec1 ^{2/-}
Onset in LD	WT	***	ns	ns	***	***	***	***
	Per1 ^{-/-}		ns	***	***	ns	ns	***
	Dec1 ^{-/-}			*	***		ns	***
	Dec2 ^{-/-}					***	***	***
	Per1 ^{-/-} Dec1 ^{-/-}					***	*	**
	Per1 ^{-/-} Dec2 ^{-/-}						ns	***
Onset variation in LD	WT	**	ns	ns	***	***	***	***
	Per1 ^{-/-}		ns	ns	*	ns	ns	***
	Dec1 ^{-/-}			ns	***		***	***
	Dec2 ^{-/-}					ns	**	***
	Per1 ^{-/-} Dec1 ^{-/-}					ns	ns	**
	Per1 ^{-/-} Dec2 ^{-/-}						ns	***
Period length in DD	WT	***	***	ns	***	***	ns	***
	Per1 ^{-/-}		***	***	ns	ns	***	***
	Dec1 ^{-/-}			***	***		ns	***
	Dec2 ^{-/-}					***	ns	***
	Per1 ^{-/-} Dec1 ^{-/-}					ns	***	***
	Per1 ^{-/-} Dec2 ^{-/-}						***	***
Period length variation in DD	WT	*	ns	ns	***	***	*	***
	Per1 ^{-/-}		ns	ns	ns	ns	ns	ns
	Dec1 ^{-/-}			ns	*		ns	***
	Dec2 ^{-/-}					***	ns	***
	Per1 ^{-/-} Dec1 ^{-/-}					ns	ns	ns
	Per1 ^{-/-} Dec2 ^{-/-}						ns	ns
Period length in LL	WT	***	ns	ns	***	***	ns	***
	Per1 ^{-/-}		***	***	ns	ns	*	***
	Dec1 ^{-/-}			ns	***		ns	***
	Dec2 ^{-/-}					***	ns	***
	Per1 ^{-/-} Dec1 ^{-/-}					ns	***	ns
	Per1 ^{-/-} Dec2 ^{-/-}						***	*
Per1 ^{-/-} Dec1 ^{2/-}						***		

ns = not significant

* p ≤ 0.05

** p ≤ 0.01

*** p ≤ 0.001

Bode et al. Table S2: Statistics of reporter gene assay

Tested parameter	Expression plasmid comparison	↓ →	Bmal1::luc, Per1	Bmal1::luc, Dec1	Bmal1::luc, Dec2	Bmal1::luc, Per1, Dec1	Bmal1::luc, Per1, Dec2	Bmal1::luc, Per1, Dec1, Dec2
Reporter gene assay	Bmal1::luc		ns	***	***	***	***	***
	Bmal1::luc, Per1			***	***	***	***	***
	Bmal1::luc, Dec1				***			***
	Bmal1::luc, Dec2					ns		***
	Bmal1::luc, Per1, Dec1						ns	***
	Bmal1::luc, Per1, Dec2						***	***

ns = not significant * $p \leq 0.05$ ** $p \leq 0.01$ *** $p \leq 0.001$

Bode et al. Table S3: Statistics of reporter gene assay

Tested parameter	Expression plasmid comparison	↓ →	Bmal1::luc, Rev-Erb alpha	Bmal1::luc, Rev-Erb alpha, Per1	Bmal1::luc, Rev-Erb alpha, Dec1	Bmal1::luc, Rev-Erb alpha, Dec2	Bmal1::luc, Rev- Erb alpha, Per1, Dec1	Bmal1::luc, Rev-Erb alpha, Per1, Dec2	Bmal1::luc, Rev-Erb alpha, Per1, Dec1, Dec2
Reporter gene assay	Bmal1::luc		***	***	***	ns	***	**	ns
	Bmal1::luc, Rev-Erb alpha			ns	***	***	**	***	***
	Bmal1::luc, Rev-Erb alpha, Per1				***	***	**	***	***
	Bmal1::luc, Rev-Erb alpha, Dec1						ns	ns	***
	Bmal1::luc, Rev-Erb alpha, Dec2				***		***	**	ns
	Bmal1::luc, Rev-Erb alpha, Per1, Dec1							***	***
Bmal1::luc, Rev-Erb alpha, Per1, Dec2								*	

ns = not significant * p ≤ 0.05 ** p ≤ 0.01 *** p ≤ 0.001