

Additional data file 6. Resampling analysis with different environmental variables.

Gene set	N genes ^a	N significant genes ^b	Empirical p value ^c	Contributing genes ^d
Core circadian	12	5	0.076 (minimum temperature)	<i>NPAS2, PER2, ARNTL2, CSNK1D, CSNK1E</i>
		0	1 (maximum temperature)	-
		2	0.1987 (short wave radiation flux)	<i>ARNTL2, CSNK1E</i>
		6	0.0191 (Δ temperature)	<i>NPAS2, PER2, ARNTL, ARNTL2, CSNK1D, CSNK1E</i>
Circadian hits (RNAi screen)	223	18	0.187 (minimum temperature)	<i>TPO, ANTXR1, CMTM7, SCHIP1, TBC1D9, PTK2, SH3GL2, ABL1, GPR158, FAM55D, BRCA2, RAB20, CTDPI, HOMER3, SLC8A2, SLC24A3, PWP2, MKL1</i>
		0	1 (maximum temperature)	-
		2	0.359 (short wave radiation flux)	<i>SCHIP1, SLC8A2</i>
		19	0.249 (Δ temperature)	<i>LZTR2, TPO, ANTXR1, CMTM7, FHIT, TBC1D9, ASB5, JAZF1, SCARA3, POPI, PTK2, SH3GL2, ABL1, GPR158, FAM55D, BRCA2, RAB20, SLC24A3, MKL1</i>
Mouse circadian/sleep disturbance	82	11	0.199 (minimum temperature)	<i>CAMK2A, EBF2, KCNB2, RORB, TNC, PRKG1, KCNMA1, NCAM1, RORA, NCOR1, APP</i>
		0	1 (maximum temperature)	-
		1	0.702 (short wave radiation flux)	<i>DRD3</i>
		12	0.192 (Δ temperature)	<i>CAMK2A, CADPS2, EBF2, RORB, TNC, PRKG1, KCNMA1, NCAM1, NOS1, RORA, NCOR1, APP</i>
Mendelian diseases causing sleep disturbance	11	4	0.195 (minimum temperature)	<i>NRXN1, HDAC4, DHCR7, SHANK3</i>

		0	1 (maximum temperature)	-
		0	1.000 (short wave radiation flux)	-
		3	0.474 (Δ temperature)	<i>NRXN1, HDAC4, SHANK3</i>
Melanopisin signaling	13	3	0.538 (minimum temperature)	<i>PRKCZ, TRPC7, PLCB4</i>
		0	1 (maximum temperature)	-
		0	1 (short wave radiation flux)	-
		4	0.285 (Δ temperature)	<i>PRKCZ, PIK3R1, TRPC7, PLCB4</i>

^a Number of genes with at least one SNP genotyped in the HGDP-CEPH panel;

^b Number of genes showing at least one significant SNP;

^c The empirical p value was calculated as described in the text and methods;

^d Genes showing at least one significant SNP.