

Supplementary Figure 1 The workflow of the study design.

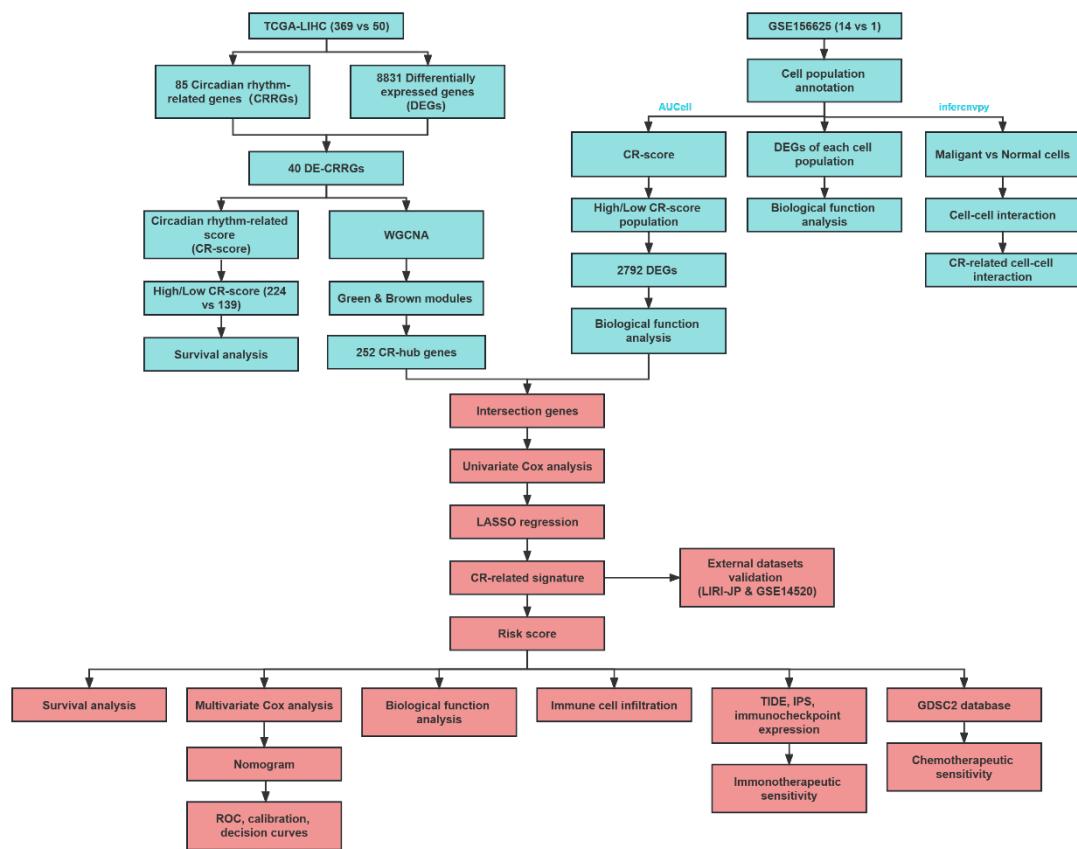


Table S1 Circadian rhythm-related genes (CRRGs) from the GeneCards.

Sources	Gene symbol
GeneCards	BMAL1
	CLOCK
	NR1D1
	RORA
	CARM1
	NCOA2
	CREBBP
	NCOA6
	PPARA
	MED1
	RXRA
	SMARCD3
	TBL1X
	TBL1XR1
	CHD9
	HELZ2
	NCOA1
	TGS1
	BHLHE40
	BHLHE41
	NPAS2
	EP300
	HDAC3
	NCOR1
	CRY1
	CRY2
	CSNK1E
	PER1
	PER2
	NAMPT
	DBP
	F7
	NOCT
	KLF15
	SERPINE1
	AVP
	BMAL2
	RAI1
	PPARGC1A
	CPT1A
	CREB1
	ATF2

CRTC2
CRTC1
SIRT1
MEF2C
MEF2D
CRTC3
SREBF1
NRIP1
PER3
LOC102724428
TPTEP2-CSNK1E
CREM
CSNK1D
SIK1
FBXL3
NR3C1
HIF1A
NFIL3
PPP1CA
PPP1CB
PPP1CC
RBM4
RPS27A
SKP1
UBA52
UBB
UBC
CUL1
BTRC
SLC46A1
LY96
HBA1
HBA2
HBB
HMOX1
APOA1
APOB
NFE2L2
CLEC1B
BACH1
TLR4
XPO1
MAFK

Table S2 DEGs between HCCs and normal samples in TCGA-LIHC cohort ($|\log_2 \text{FC}| > 0.585$ & $P < 0.05$).

Total	Up	Down
8831	7793	1038

Table S3 Venn analysis of the 40 intersected CRRGs between CRRGs from GeneCards and DEGs from TCGA-LIHC cohort.

Intersected CRRGs	Gene symbol
	CLEC1B
	CRTC2
	SKP1
	CSNK1D
	PPP1CA
	PPP1CC
	MEF2D
	XPO1
	CSNK1E
	SLC46A1
	HDAC3
	UBA52
	RPS27A
	CRTC1
	NCOA6
	HBB
	RBM4
	DBP
	HBA2
	TGS1
	TBL1XR1
	ATF2
	MED1
	PPP1CB
	SERPINE1
	NPAS2
	NOCT
	NCOA2
	CARM1
	TLR4
	SMARCD3
	NFIL3
	RAI1
	HMOX1
	NAMPT
	APOA1
	BHLHE40
	BMAL2
	PPARGC1A
	LY96

Table S4 Identification of the hub genes based on WGCNA.

Hub genes	Gene symbol
	ABI3BP
	ABR
	AC006062.1
	ACTA2
	ADAMTS1
	ADAMTS2
	ADAMTS4
	ADCY3
	ADGRA2
	ADGRF5
	AEBP1
	AKT3
	ANTXR1
	AOC3
	AP001189.1
	AP001189.3
	ARAP3
	ARHGAP23
	ARHGAP31
	ARHGAP6
	ASPN
	BGN
	BX322234.1
	C7
	CACNA1C
	CALCRL
	CALHM2
	CCDC80
	CD248
	CD93
	CDH11
	CDH5
	CDR2L
	COL12A1
	COL14A1
	COL15A1
	COL16A1
	COL1A1
	COL1A2
	COL3A1
	COL4A1
	COL4A2

COL5A1
COL6A1
COL6A2
COL6A3
COL8A1
CPXM2
CRISPLD2
CTTNBP2NL
CXCL12
CYTH3
DACT3
DCN
DENND3
DHH
DKK2
DSE
ECM1
EDNRA
EFEMP1
ELK3
EMILIN1
EMP1
ENDOD1
ENG
EPHA3
ERG
ETS1
F2R
F3
FAP
FBLN2
FBLN5
FBN1
FIBIN
FILIP1L
FLNA
FLT4
FMOD
FNDC1
FOXF1
FSTL1
GAS1
GEM
GFPT2

GGT5
GIMAP6
GPR4
GUCY1A1
GUCY1A2
GUCY1B1
HAND2
HAND2-AS1
HCG11
HDGFL3
HECW2
HEG1
HEPH
HEYL
HSPG2
ID4
IGFBP5
IGFBP7
IQGAP1
IRAG1
ISLR
ITGA11
ITGA3
ITGA9
JAM3
KCNE4
KCTD12
KIF26A
KIRREL1
LAMA2
LBH
LDB2
LHFPL6
LINC00924
LINC01197
LINC01679
LRRC32
LTBP2
LTBP4
LUM
LXN
LZTS1
MAP3K6
MEDAG

MFAP4
MMP2
MOXD1
MRC2
MSRB3
MXRA5
MYCT1
MYH11
MYOF
NAV1
NEXN
NFASC
NOS3
NOTCH3
NOTCH4
OMD
OSBPL5
P4HA3
PCDH17
PCDHGB7
PCDHGC3
PDE1A
PDGFRA
PEAR1
PFKFB3
PHLDB1
PLAT
PLEKHG1
PLEKHG5
PLN
PLXDC2
PMEPA1
PODN
PODXL
POSTN
PPP1R12B
PPP1R18
PRELP
PREX1
PRKG1
PRSS23
PTGIR
PTGIS
PTPRB

PTPRE
PXDN
RAB31
RBMS2
RCAN2
SCARF1
SLC6A6
SLIT2
SLIT3
SMOC2
SNAI1
SPARC
SPON1
SPRY1
SRPX
SSC5D
ST8SIA6
STING1
SULF1
SVEP1
SYNPO2
SYT11
TAGLN
TANC1
TBXA2R
TCF21
TCF4
TEK
TGFB3
THBD
THBS2
TIMP2
TMEM255B
TNC
TNXB
TPM4
TSHZ3
UNC5B
VCAN
VCL
VEGFC
ZEB2
ZFPM2
ZSWIM4

AC000089.1
AC004453.1
AC034236.1
AC091153.1
AC125807.1
AL009174.1
EEF1B2P6
FBL
RPL10AP6
RPL11
RPL13
RPL14
RPL14P1
RPL24
RPL24P2
RPL27
RPL27A
RPL29
RPL32
RPL34
RPL37
RPL37A
RPL37P6
RPL38
RPL7AP50
RPLP2
RPS14
RPS15A
RPS21
RPS24
RPS24P8
RPS25
RPS29
RPS7

Table S5 DEGs between high CR-score and low CR-score groups in GSE156625 dataset ($|\log_2 \text{FC}| > 0.585$ & $P- < 0.05$).

Total	Up	Down
2792	2649	143

Table S6 Venn analysis of the 44 intersected CRRCs between CRRGs from WGCNA and scRNA-seq data analysis.

Intersected CRRGs	Gene symbol
	RPS29
	IQGAP1
	RPS25
	TAGLN
	RPLP2
	RPL27A
	PTPRE
	CALHM2
	PLXDC2
	ETS1
	PFKFB3
	ACTA2
	VCL
	RPS24
	TPM4
	ZSWIM4
	PMEPA1
	PREX1
	RPS21
	RPL13
	CRISPLD2
	MYH11
	RPS15A
	RAB31
	RPL27
	RPL38
	RPL37A
	RPL32
	SLC6A6
	RPL14
	ZEB2
	RPL34
	RPL24
	RPL29
	RPL11
	RPS7

LBH
SYT11
FLNA
GEM
RPS14
RPL37
DSE
PPP1R18
