Supplementary information

Inhibition of JNK/c-Jun-ATF2 Overcomes Cisplatin Resistance in Liver Cancer through down-

Regulating Galectin-1

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Figure S1. CDDP activated JNK signal in liver cancer cells. IC_{50} of CDDP was performed in Hep 3B, Hep 3B/DR (A) and SK-Hep 1 cells (B). (C) Dual-reporter assay detected the activity of ATF element. The levels of firefly luciferase activity were normalized to Renilla luciferase activity, which reflected the relative activity of ATF element. **P < 0.01, ***P < 0.001.



Figure S2. Inhibition of JNK/c-Jun-ATF2 by SP600125 or ATF-decoy decreased the activity of ATF element in liver cancer cells. (A) The relative activity of ATF elements was detected by dual-luciferase reporter assay after SP600125 and CDDP treatment. (20 μ M CDDP, 20 μ M SP600125 for 18 h) **P* < 0.05, ***P* < 0.01, ****P* < 0.001. (B) Cells were co-transfected with ATF-decoy or mut-ATF-decoy (mut) and pGL3-ATF-Flue plasmids. The dual-reporter analysis was performed. **P* < 0.05, ***P* < 0.01, ****P* < 0.001.



Figure S3. Inhibition of JNK/c-Jun-ATF2 by SP600125 or ATF-decoy enhanced DNA damage in liver cancer cells. (A, B) γ -H2AX was analyzed by IF in Hep 3B and SK-Hep 1 cells. The representative pictures were shown. The nuclei were stained with DAPI. Scale bar, 10 µm. The quantification of the percentage of foci positive cells was shown in Figure 2E, F. (C, D) Comet assay was performed for detecting DNA damage in Hep 3B and SK-Hep 1 cells. The representative pictures were shown. Scale bar, 100 µm. The quantification of the average tail moment per cell was shown in Figure 2G, H.



Figure S4. C2/c-Jun and C2/ATF2 enhanced the transcription of *Galectin-1* in liver cancer cells. (A, B) Cells were transfected with C2/c-Jun and C2/ATF2. The activity of Gal-1-WT-Fluc or Gal-1-mut-Fluc was measured by dual-reporter assay. (C) The *Galectin-1* mRNA levels were detected by qPCR. *P < 0.05, **P < 0.01, ***P < 0.001.



Figure S5. Inhibition of JNK/c-Jun-ATF2 by SP600125 or ATF-decoy decreased the mRNA level of *Galectin-1* in liver cancer cells. (A) Liver cells were treated with SP600125 and CDDP (20 μ M CDDP, 20 μ M SP600125 for 18 h). The *Galectin-1* mRNA levels were detected by qPCR. (B) Liver cancer cells were transfected with ATF-decoy or mut-ATF-decoy (mut) and treated with CDDP (20 μ M for 18 h). The *Galectin-1* mRNA levels were detected by qPCR. (B) Liver cancer cells were I mRNA levels were detected by qPCR. (P = 0.05, **P < 0.01.



Figure S6. Inhibition of JNK/c-Jun-ATF2 by SP600125 or ATF-decoy decreased the protein expression of Galectin-1 in liver cancer cells. (A, B) The expression of Galectin-1 in western blot was quantitated by the gray density analysis. The representative images were shown in Figure 3J and K. *P < 0.05, **P < 0.01.



Figure S7. Galectin-1 was knocked down by shRNA. Liver cancer cells were transfected with sh Galectin-1. Galectin-1 protein was detected by western blot. A control shRNA and four Galectin-1-target shRNA were designed, represented as NC, 1, 2, 3 and 4 respectively.



Figure S8. *Galectin-1* knockdown reversed CDDP resistance in liver cancer cells. (A, B) Cells with *Galectin-1* knockdown were treated with CDDP (20 μ M for 18 h). The expression of γ -H2AX and cleaved caspase 3 was quantitated by the gray density analysis. The representative images were shown in Figure 4A. *P < 0.05, **P < 0.01.



Figure S9. Galectin-1 hindered the efficacy of CDDP by mediating DNA homologous recombination repair. (A) Liver cancer cells were transfected with sh Galectin-1. γ H2AX was analyzed by immunofluorescence. Cells were stained by γ -H2AX primary antibody and DAPI. The corresponding quantification results refer to figure 5B. Scale bar, 10 µm. (B) Liver cancer cells were transfected with sh Galectin-1. Comet assay was performed for detecting DNA damage. The corresponding quantification results refer to figure 5C. Scale bar, 100 µm. (C) The principle of determination of HR or NHEJ frequency. (D) Quantification of HR or NHEJ frequency in Galectin-1-overexpressing HEK293T cells.



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Figure S10. High expression of Galectin-1 is related to poor prognosis in liver cancer. (A) Expression profile of *Galectin-1* in liver cancer versus control tissues in TCGA dataset. (B) The overall survival analysis between high and low *Galectin-1* expression of liver cancer.

Supplementary Tables

Table S1

Name	Company	Catalogue
DMEM	Gibco	C11995500BT
RPMI-1640	Gibco	C11875500BT
fetal bovine serum	Gibco	10270-106
Cisplatin	Selleck Chemicals	S1166
SP600125	Selleck Chemicals	S1460
puromycin	Meilunbio	MA0318
BCA assay	Beyotime	P0010
Human Phospho-Kinase Array	R&D	ARY003C
RIPA	Beyotime	P0013C
protease inhibitor cocktail	Thermo Fisher	78430
loading buffer	EpiZyme	LT101S
PVDF membranes	BioTrace	66485
Dynabeads [®] Protein A	Invitrogen	1001D
Dual-Luciferase® Reporter Assay	Promega	P1910
BSA	Aladdin	A104912
Dylight 549-conjugated goat anti-rabbit IgG	Abbkine	A23320
DAPI	Solarbio	S2110
Alkaline comet	Trevigen	4250-050-К
SYBR Green I	Solarbio	SY1020
Total RNA Kit I	Omega	R6834-02
PrimeScript TM RT reagent Kit	Takara	RR047A
TB Green [®] Premix Ex TaqTM II	Takara	RR820A
ViviRen TM	Promega	P1232
_D -luciferin	PerkinElmer	122799
Galacto-Star TM System	Thermo Fisher	T1012
MaxVision TM HRP-Polymer anti-Rabbit IHC Kit	MXB Biotechnologies	KIT-5005
DAB Kit (20×)	MXB Biotechnologies	DAB-1031

Table S2

Antibody	Company	Catalogue	WB/IP/IHC/IF/ChIP	dilution
JNK	CST	9252S	WB	1:1000
c-Jun	CST	9165S	WB/IP/ChIP	1:1000/1:50/1:50
c-Jun	CST	23158	WB	1:1000
ATF2	CST	35031S	WB/IP/ChIP	1:1000/1:50/1:50
p-JNK	CST	4668S	WB	1:1000
p-c-Jun	CST	32708	WB/IHC	1:1000/1:200
p-ATF2	CST	92258	WB/IHC	1:1000/1:200
γ-H2AX	CST	9718S	WB/IF	1:1000/1:200
Galectin-1	CST	129368	WB	1:1000
Galectin-1	CST	13888	IHC	1:250
GAPDH	Proteintech	60004-1-Ig	WB	1:5000
rabbit IgG	CST	2729	ChIP	2 μL

Table S3

Name	Sequence	
ATF-decoy	AATTC <u>TTACCTCA</u> GATGA <u>TTACCTCA</u> TCCCGC <u>TTA-</u> <u>CCTCA</u> CGATCAG <u>TTACCTCA</u> C	
mut-ATF-decoy	AATTC <u>GGACCTCG</u> GATGA <u>GGACCTCG</u> TCCCGC <u>GGA-</u> <u>CCTCG</u> CGATCAG <u>GGACCTCG</u> C	
2×ATF	TCCCGC <u>TTACCTCA</u> CGATCAG <u>TTACCTCA</u> C	
<i>MYL6B</i> primers	forward 5'-GGAGCCTCCAGTCGATCTCT-3' reverse 5'-GCAGGAAAGTCTCAAAGTCCAC-3'	
<i>RHOC</i> primers	forward 5'-AGACCTGCCTCCTCATCGT-3' reverse 5'-GGCCGCAGTCGATCATAGTC-3'	
HCFC primers	forward 5'-ATCGAGTCCCTGGGTGTGA-3' reverse 5'-GGTGCCCAAATCATCGTCTG-3'	
Galectin-1 primers	forward 5'-TCGCCAGCAACCTGAATCTC-3' reverse 5'-GCACGAAGCTCTTAGCGTCA-3'	
MAST2 primers	forward 5'-ACGGAGGTGGTAGAGCTGAT-3' reverse 5'-TAAGGGAAGAAAGGCTGCGG-3'	
ACTB primers	forward 5'-GAGAAAATCTGGCACCACACC-3' reverse 5'-GGATAGCACAGCCTGGATAGCAA-3'	
Primers for ChIP-qPCR	P-qPCR forward 5'-TCAGAGGAGATGTTAAGAGAGCAGAC- reverse 5'-CGTTGGCCAGGCTGGTC-3'	

Supplementary methods

1. CCK-8 assay IC₅₀ of CDDP.

For the determination of IC_{50} of CDDP in liver cancer cells, about 7000 cells/well were cultured in the 96well plates. The cells were treated with CDDP at different concentrations for 24 h. CCK-8 reagent was diluted in the culture medium (10%). Each well was added 100 μ L CCK-8 working solution and incubated in 37 °C incubators for 1.5 h. The absorbance of 450 nm was detected using Multimode Reader Synergy HTX (BIO-RAD).

2. Plasmids.

Plasmids pGL3-2×ATF-Fluc, pcDNA3.1-C2/c-Jun and pCMV-C2/ATF2 were gifts from Professor Mingtao Li (Zhongshan School of Medicine, Sun Yat-sen University). pLR-Renilla Luciferase (Rluc) was purchased from Promega.

3. qPCR analysis

Cells were transfected with ATF-decoy, mut-ATF-decoy, pcDNA3.1-C2/c-Jun or pCMV-C2/ATF2, and then treated or not treated with CDDP. RNA was obtained for qPCR assay. The relative expression of *Galectin-1* was analyzed according to the $\Delta\Delta$ CT relative quantification method and normalized to the expression of *ACTB*. Expression measurements of mRNA were performed in triplicate. The kit information was shown in Table S1. The primers are listed in Table S3.

4. Dual-luciferase reporter assay

pGL3-2×ATF-Fluc and pLR-Renilla plasmid were co-transfected to cells at ratio of 9:1 before drug treatment. To detect the effect of declined c-Jun-ATF2 on ATF element, cells were also co-transfected with the two plasmids above and ATF-decoy or mut-ATF-decoy. To determine the function of C2/c-Jun and C2/ATF2 on *Galectin-1* prompter, cells were co-transfected with Gal-1-WT-Fluc (or Gal-1-mut-Fluc) and pcDNA3.1-C2/c-Jun and pCMV-C2/ATF2. Dual-Luciferase® Reporter Assay was used to detect the activity of firefly and renilla luciferase.

5. Determination of HR or NHEJ frequency.

The plasmid system was a gift from Professor Yong Zhao (School of Life Sciences, Sun Yat-sen University, Guangzhou, China). This system was used to detect the frequency of homologous recombination (HR) or non-homologous end-joining (NHEJ) repair. The analysis was performed as the published paper [1, 2]. It consists

of three plasmids, including the DSB reporter plasmid, DSB repair templet plasmid and *LacZ* plasmid (transfection control). The three plasmids and pcDNA-Galectin-1 were co-transfected to HEK293T cells for 72 h. The luciferase activity was detected by Dual-Luciferase® Reporter Assay. The LacZ activity was detected by Galacto-StarTM System. The activity of firefly or renilla luciferase was normalized to LacZ activity.

6. Galectin-1 expression and survival analysis of liver cancer based on TCGA dataset.

The expression of *Galectin-1* mRNA in liver cancer was obtained from TCGA. The expression analysis and survival analysis were performed at http://ualcan.path.uab.edu/ and http://gepia2.cancer-pku.cn/#index.

- Zhang C, Chen L, Peng D, et al. METTL3 and N6-Methyladenosine Promote Homologous Recombination-Mediated Repair of DSBs by Modulating DNA-RNA Hybrid Accumulation. Mol Cell. 2020; 79: 425-42 e7.
- Xie C, Chen YL, Wang DF, et al. SgRNA Expression of CRIPSR-Cas9 System Based on MiRNA Polycistrons as a Versatile Tool to Manipulate Multiple and Tissue-Specific Genome Editing. Sci Rep. 2017; 7: 5795.