

# CAFs-Derived Exosomal miRNA-130a Confers Cisplatin Resistance of NSCLC Cells Through PUM2-Dependent Packaging

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**Purpose:** Chemoresistance is a significant barrier to the treatment and management of non-small cell lung cancer (NSCLC). Exosomes play an essential role in intercellular communication. Understanding the mechanism underlying the role of tumor stroma, especially cancer-associated fibroblasts (CAFs), during chemoresistance would significantly contribute to the clinical application of chemotherapy agents.

**Results:** In this study, we demonstrated that NSCLC-derived CAFs were innately resistant to cisplatin treatment and CAFs-conditioned medium significantly promoted the survival rate of NSCLC cells after cisplatin treatment. Additionally, CAFs-derived exosomes were taken up by NSCLC cells. Moreover, exosomal miRNA-130a was transferred from CAFs to recipient NSCLC cells and knockdown of miRNA-130a reversed the effect of CAFs-derived exosomes during chemoresistance of NSCLC cells. Furthermore, pumilio homolog 2 (PUM2), a RNA-binding protein, mediated the packaging of miRNA-130a into exosomes. The overexpression and knockdown of PUM2 promoted and inhibited tumor growth of xenograft mice, respectively.

**Conclusion:** Taken together, these results suggest that CAFs-derived exosomes confer cisplatin resistance of NSCLC cells through transferring miRNA-130a and that PUM2 is a critical factor for packaging miRNA-130a into exosomes. This study indicates that CAFs-derived exosomal miRNA-130a may be a potential therapeutic target for cisplatin resistance in NSCLC.

**Keywords:** chemoresistance, non-small cell lung cancer, cancer-associated fibroblasts, exosome, miRNA-130a, pumilio homolog 2

## Introduction

Lung cancer has emerged as the leading cause of cancer-associated deaths worldwide, accounting for about 20% of all cancer deaths.<sup>1</sup> In 2018, approximately 2.09 million new lung cancer cases took place, ranking first in all cancer types.<sup>2,3</sup> In China, lung cancer mortality is higher than in most other countries, and it is estimated that lung cancer mortality may surge nearly 40% between 2015 and 2030.<sup>4</sup> As the primary type of lung cancer, non-small cell lung cancer (NSCLC) exhibits multiple histological types, such as bronchoalveolar carcinoma, large cell carcinoma, squamous cell carcinoma, and adenocarcinoma.<sup>5</sup> To date, platinum-based chemotherapy is the most effective therapy to treat advanced NSCLC patients,<sup>6</sup> with a significantly higher survival and response rate.<sup>7,8</sup> Alternatively, cisplatin (cis-Diamminedichloro-platinum(II))-based therapeutic strategy has

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a relatively low response rate, significant side effects, and acquired or intrinsic chemoresistance.<sup>9,10</sup> As such, it is urgent to improve the efficacy of chemotherapy to NSCLC and explore the molecular mechanism underlying cisplatin resistance of NSCLC.

Cancer-associated fibroblasts (CAFs) have been demonstrated to be essential stromal components in the tumor microenvironment, regulating cancer cell growth, migration, invasion, and chemo-response through various mechanisms.<sup>11</sup> Chemoresistance, either acquired or intrinsic, is associated with a complicated and multifactorial process, including decreases of intracellular drugs, changes of drug targets, aberrant regulation of cell survival or death, and crosstalks between tumor microenvironment and cancer cells.<sup>12,13</sup> Growing evidence recently reveals that CAFs play a critical role in the regulation of chemoresistance in various cancer types. For example, CAFs can secrete cytokines or growth factors to promote tumorigenesis and chemoresistance in breast cancer.<sup>14</sup> In addition, CAFs-derived interleukin 6 (IL6) is essential for chemoresistance through increasing C-X-C motif chemokine receptor 7 (CXCR7) expression.<sup>15</sup> Moreover, increased secretion of epidermal growth factor sufficiently insensitizes neighboring tumor cells to cetuximab treatment via the mitogen-activated protein kinases (MAPK) pathway.<sup>16</sup> Furthermore, CAFs have been reported to promote cisplatin resistance in bladder cancer cells through activating the IGF-1/ERβ/Bcl-2 signaling pathway.<sup>17</sup>

Given the importance of CAFs in chemoresistance of cancers, the mechanism underlying how CAFs influence the cellular activities, especially the communication between CAFs and cancer cells, has drawn considerable attention from both laboratory and clinical research fields. Over the past decade, it has been demonstrated that exosomes are essential molecule/information transporters in intercellular communication, including the interaction between tumor cells and CAFs.<sup>18,19</sup> Exosomes, membrane-derived microvesicles (30–150 nm), are secreted by almost all cell types and are extensively distributed in various body fluids.<sup>20</sup> Exosomes can transport bioactive components, such as proteins, non-coding RNAs, DNA fragments, mRNAs, and lipids, from donor cells to neighboring cells, thereby impacting the physiological activities of recipient cells.<sup>21,22</sup> Among these functional components, microRNAs (miRNAs), a class of approximately 22-nucleotide non-coding RNAs, have been extensively reported to play a critical role in exosome-mediated cell-to-cell communication.<sup>23,24</sup> For example, CAFs-derived exosomal miRNA-106b can promote gemcitabine resistance of pancreatic cancer cells through silencing

TP53INP1.<sup>25</sup> In addition, exosomal miRNA-522 secreted from CAFs enhances acquired chemoresistance to cisplatin in gastric cancer.<sup>26</sup> In addition, exosomal miRNA-21 derived from CAFs contribute to oxaliplatin resistance in colorectal cancer.<sup>27</sup>

To the best of our knowledge, the role of CAFs-derived exosomal miRNAs during chemoresistance of NSCLC cells has not been thoroughly investigated. Therefore, this study aimed to determine the effect of CAFs-derived exosomes during the chemoresistance of NSCLC cells. In this study, we demonstrated that CAFs-derived exosomes significantly improved the survival rate of NSCLC cells in response to cisplatin treatment. In addition, the experiments suggest that miRNA-130a is a functional miRNA which mediates the role of CAFs-derived exosomes and that pumilio homolog 2 (PUM2) may be a key factor for packaging miRNA-130a into exosomes.

## Materials and Methods

### Ethics Statement

In this study, written informed consent was offered by the participants before enrollment. The experimental protocols were approved by the Ethics Committee of Linyi chest Hospital and were conducted based on the Declaration of Helsinki. All animal studies were approved by the Institute Animal Care and Use Committee of Linyi chest Hospital and were performed according to the Guide for the Care and Use of Laboratory Animals.<sup>28</sup> The mice used in this study were raised in the medicine animal facilities of Linyi chest Hospital.

### Samples and Patients

All clinical samples were collected from Linyi chest Hospital. Twenty pairs of tumor and adjacent normal tissues were obtained from twenty patients who were diagnosed with NSCLC based on the evidence of pathological histology from 2016–2018. [Table 1](#) shows the patient information. None of these patients underwent chemotherapy and radiotherapy before the surgery. The medical records and follow-up data of these NSCLC patients were retrospectively reviewed, and pathological and clinical characteristics were also respectively reviewed.

### Cell Culture

The NSCLC cell lines, A549 and HCC827, were obtained from the Cell Bank of China Academy of Sciences

**Table 1** Patient Characteristics

Variable	Patients
Patients	20
Gender	
Male (%)	9 (45)
Female (%)	11 (55)
Mean age (range)	57 (24–71)
Histology	
Adenocarcinoma (%)	12 (60.0)
Squamous (%)	8 (40.0)
Differentiation	
Poor (%)	16 (80.0)
Moderate (%)	3 (15.0)
Well (%)	1 (5.0)
Metastatic organ	
≥3 (%)	9 (45.0)
<3 (%)	11 (55.0)
Smoking history	
Never (%)	2 (10.0)
Former (%)	8 (40.0)
Current (%)	7 (35.0)
Unknown (%)	3 (15.0)

(Shanghai, China). To establish the cisplatin-resistant cell line, A549-R, A549 cells were treated with a stepwise increasing concentration of cisplatin (Sigma-Aldrich, USA) as previously described.<sup>29,30</sup> The CAFs and normal fibroblasts (NFs) were isolated from NSCLC tissues and adjacent normal tissues by primary culture, as previously described, respectively.<sup>31,32</sup> Both third-generation purified CAFs and NFs were further identified by detecting CAFs-specific markers, including  $\alpha$ -smooth muscle actin ( $\alpha$ -SMA; Thermo, USA), fibroblast activation protein (FAP; Thermo, USA), and ferroptosis suppressor protein 1 (FSP1; Abcam, China). The A549 and HCC827 cells were cultured in RPMI 1640 medium (GIBCO-BRL, USA) supplemented with 10% fetal bovine serum (FBS; GIBCO-BRL, USA), 100 U/mL penicillin, and 100  $\mu$ g/mL streptomycin in humidified air with 5% CO<sub>2</sub> at 37 °C. The coculture system was created using transwell membranes with 0.4  $\mu$ m pores (Corning, USA) in a 6-well plate. The CAFs ( $1 \times 10^5$ ) were treated with 20  $\mu$ M GW4869 (Sigma-Aldrich, China) for 24 hours and placed above the membranes. The A549 and HCC827 cells were placed below the permeable membranes. The cells were placed in the coculture system for 72 hours in humidified air with 5% CO<sub>2</sub> at 37 °C.

## Western Blots

Exosomes or cells were collected at specific times, and total proteins were isolated using SDS lysis buffer (Sigma-Aldrich, China). Total proteins (20  $\mu$ g) were separated by polyacrylamide gels (15%/10%) and then transferred onto PVDF membranes (Merck Millipore, USA). The blots were blocked with 5% bovine serum albumin (BSA; Sigma-Aldrich, China) at room temperature for 1 hour and incubated with primary antibodies in a microplate shaker overnight at 4 °C. The expression of  $\beta$ -actin was used throughout as a loading control, and all other primary antibodies were used according to the manufacturer's instructions. Next, the membranes were incubated with corresponding secondary antibodies and the protein bands were detected using the Bio-Rad Chemidoc system (Bio-Rad Laboratories, USA).

## Real-Time PCR

Total RNAs were isolated from cells, exosomes, and tissues using the TRIzol reagent (Invitrogen, USA), and first-strand cDNAs were synthesized using the Prime Script RT reagent Kit (Takara, Japan) according to the manufacturer's instructions. The miRNA cDNAs were synthesized from total RNAs of cells or tissues using the miRNA First-Strand cDNA Synthesis Kit (TIANGEN, China). Real-time PCR for mRNAs was performed using the SYBR Premix Ex Taq Reagent Kit (Takara, Japan) on a 7500 Real-Time PCR Platform (Applied Biosystems, USA). Real-time PCR for miRNA was performed using the miRcute Plus miRNA qPCR Detection Kit (TIANGEN, China).  $\beta$ -actin and U6 were used as internal controls for analyzing relative expressions of RNAs and miRNAs, respectively. The  $2^{-\Delta\Delta Ct}$  method was carried out to analyze relative expressions.<sup>33</sup> Table 2 summarizes the primer information.

## MiRNA Array

The NFs- and CAFs-derived exosomes were used to determine the miRNA expression profile. Exosomal RNAs were isolated as the method mentioned above. The purification and concentration of RNAs were determined by NanoDrop ND-2000 (Thermo Scientific, USA). Exosomal RNA was labeled using FlashTag™ Biotin HSR RNA Labeling Kits (Thermo Scientific, USA) followed by the hybridization of labeled exosomal RNA. After hybridization, the array was scanned and the data was processed. Analysis for miRNA expression profile was performed using the 7G platform Affymetrix

**Table 2** Primer Information

Gene	Primer Sequences
MRP2	F: 5'-CCCTGCTGTTTCGATATACCAATC-3' R: 5'-TCGAGAGAATCCAGAATAGGGAC-3'
ATP7B	F: 5'-ATATTGAGCGGTTACAAAGCACT-3' R: 5'-TGCCCCAAGGTCTCAGAATTA-3'
CTR1	F: 5'-GGGGATGAGCTATATGGACTCC-3' R: 5'-TCACCAAACCGAAAAACAGTAG-3'
XIAP	F: 5'-AATAGTGCCACGCAGTCTACA-3' R: 5'-CAGATGGCCTGTCTAAGGCAA-3'
ERCC1	F: 5'-CTACGCCGAATATGCCATCTC-3' R: 5'-GTACGGGATTGCCCTCTG-3'
ERCC4	F: 5'-CCTCTTTCGCCAGAAAAACAAAC-3' R: 5'-TTTACTGCTACATGGAACCTTGG-3'
HRS	F: 5'-AGTGGCTGTCGGGTATTCATC-3' R: 5'-CCGTCCATATCCCTTGAAGAATC-3'
TSG101	F: 5'-GAGAGCCAGCTCAAGAAAATGG-3' R: 5'-TGAGGTTCAATAGTTCCTGGA-3'
STAM1	F: 5'-AATCCCTTCGATCAGGATGTTGA-3' R: 5'-CGAGACTGACCAACTTTATCACA-3'
VPS4B	F: 5'-ATGTCATCCACTTCGCCAAC-3' R: 5'-TTGCTTGGCTTTATCACCTG-3'
ALIX	F: 5'-GGTGCAGCTGAAGAAGACCT-3' R: 5'-CAGGTTCTGCTCTGCAAT-3'
CD9	F: 5'-TCCACTATGCGTTGAACTGCT-3' R: 5'-GGTTTCGAGTACGTCCTTCTTG-3'
CD63	F: 5'-ATGCAGGCAGATTTTAAGTGCT-3' R: 5'-GTTCTTCGACATGGAAGGGATT-3'
nSMase2	F: 5'-GCTGCCCTTTGCGTTTTCTC-3' R: 5'-TCCAGCCGTGAATAGATGTAGG-3'
PLD2	F: 5'-CAGATGGAGTCCGATGAGGTG-3' R: 5'-CCGCTGGTATATCTTTCGGTG-3'
DGKa	F: 5'-AAGAAGCTTCGTGCATGCAGGCC-3' R: 5'-GGAGGATCCGCGAAAACATACCACCTATC-3'
RAB11A	F: 5'-CAACAAGAAGCATCCAGTTGA-3' R: 5'-GCACCTACAGCTCCACGATAAT-3'
RAB35	F: 5'-TACTGTTGCGTTTTGCAGACA-3' R: 5'-CCCCGATAATACGTGGAGGTG-3'
RAB2B	F: 5'-GTACGACATTACAAGGCGTGA-3' R: 5'-ATGTTGGAAGTAGAGTGCTGC-3'
RAB5A	F: 5'-CAAGGCCGACCTAGCAAATAA-3' R: 5'-GATGTTTTAGCGGATGTCTCCAT-3'

(Continued)

**Table 2** (Continued).

Gene	Primer Sequences
RAB9A	F: 5'-AGGGACAACGGCGACTATC-3' R: 5'-TCTGACCTATCCTCGGTAGCA-3'
RAB27A	F: 5'-GCTTTGGGAGACTCTGGTGA-3' R: 5'-TCAATGCCCACTGTTGTGATAAA-3'
RAB27B	F: 5'-TAGACTTTTCGGGAAAAACGTGTG-3' R: 5'-AGAAGCTCTGTTGACTGGTGA-3'
RAB7	F: 5'-GTGTTGCTGAAGGTTATCATCCT-3' R: 5'-GCTCCTATTGTGGCTTTGTACTG-3'
GSH	F: 5'-AGATGTTCAATGAGCTGCTCACG-3' R: 5'-CCCAGTAGAAGTCTGCCAAGTTAC-3'
YKT6	F: 5'-TGGTCACTCAGTAGATACCAG-3' R: 5'-CTCGCTTAACAGAGACTCCA-3'
PKM2	F: 5'-ATAACGCCTACATGGAAAAGTGT-3' R: 5'-TAAGCCCATCATCCACGTAGA-3'
ATG3	F: 5'-ATGATCCCTGTAACCTTAGCCCA-3' R: 5'-CACGGAAGCAAACAACCTCAAC-3'
ATG7	F: 5'-ATGATCCCTGTAACCTTAGCCCA-3' R: 5'-CACGGAAGCAAACAACCTCAAC-3'
$\beta$ -actin	F: 5'-TCACCCACACTGTGCCATCTACGA-3' R: 5'-CAGCGGAACCGCTCATTGCCAATGG-3'
miRNA-130a	F: 5'-GTCAGTGCAATGTTAAAGGGCAT-3' R: 5'-CAGTGCCTGTCGTGGAGT-3'
U6	F: 5'-CTCGCTTCGGCAGCACA-3' R: 5'-AACGCTTCACGAATTTGCGT-3'

GeneChip Command Console Software (version 4.0). The miRNA expression profile was shown in Volcano Plotting. The miRNAs had  $\geq 2$ -fold change and adjusted  $P < 0.05$  were considered as differentially expressed miRNAs.

## Immunofluorescence Assay

Cells ( $1 \times 10^5$ ) cultured on cover slips were fixed with 4% paraformaldehyde (Thermo Fisher Scientific, China), permeabilized with 0.1% Triton X-100 (Abcam, China), and then blocked in 3% BSA (Sigma-Aldrich, China). Thereafter, processed cells were incubated with primary antibodies at 4 °C overnight and then incubated with corresponding secondary antibodies for 30 mins at room temperature in the dark. Subsequently, samples were treated with 4',6-diamidino-2-phenylindole (DAPI; Invitrogen, USA) to label the nuclei. Cells were imaged using a SP8 laser-scanning confocal microscope (Leica Microsystems, Germany).



## MTT Assay

Cell viability was determined using the MTT assay. Cells were administered a treatment as indicated and then seeded in a 96-well plate (5000 cells per well) in triplicate. Twelve hours later, cells were treated with different concentrations of cisplatin for 72 hours. Next, the viability of cells was determined using the MTT assay kit (Abcam, USA) according to the manufacturer's instructions. Absorbance was read at 490 nm. Meanwhile, cell proliferation was determined by seeding cells in a 96-well plate (2000 cells per well) in triplicate after pretreatment as indicated. Cell growth was determined daily by recording the absorbance at 490 nm in a plate reader (Molecular Devices, USA).

## Conditioned Medium

Cells ( $1 \times 10^5$ ) were seeded in a culture dish for 24 hours and then the culture medium was replaced with serum-free DMEM (GIBCO-BRL, USA) and incubated for 48 hours. Conditioned medium (CM) with exosomes was prepared by removing exosomes through successively centrifuging at  $300 \times g$  for 20 mins,  $2000 \times g$  for 20 mins, and  $12,000 \times g$  for 70 mins. Cisplatin-treated cells CM was prepared by culturing cells in serum-free DMEM supplemented with  $10 \mu\text{M}$  cisplatin.

## Exosome Isolation and Identification

Exosomes were isolated from CM with different pretreatments through successively centrifuging steps.<sup>34,35</sup> The morphology and size distribution of exosomes were determined by FEI Tecnai G2 Spirit transmission electron microscopy (Thermo Scientific, USA) and nanoparticle tracking analysis (NTA), respectively, as previously described.<sup>36,37</sup> Exosomal protein was determined using the BCA Protein Assay Kit (Abcam, China).

## Exosomes Immunofluorescence

Exosomes were labeled with the lipophilic dye, DiO (Biotium, USA), according to the manufacturer's instructions. The A549 cells ( $1 \times 10^5$ ) seeded on cover slips were cocultured with labeled exosomes for 24 hours in an 8-well plate. The CAFs with Cy3-labeled miRNA-130a (Biocompare, USA) were cocultured with A549 cells for 48 hours in an 8-well plate. The cytoskeleton of A549 cells was labeled with TRITC Phalloidin or FITC Phalloidin (Sigma-Aldrich, China) according to the manufacturer's instructions. Cells were then used for the immunofluorescence assay as mentioned above.

## Cell Transfection

The cDNA of PUM2 and EIF4B (with 3' UTR) was synthesized using PrimeSTAR™ HS DNA Polymerase (Takara, Japan) and then were inserted into the PGMLV-6395 vector (Genomeditech, China). The miRNA-130a mimics, miRNA-130a mimics negative control (miRNA-130a mimics-NC), miRNA-130a inhibitor, miRNA-130a inhibitor negative control (miRNA-130a inhibitor-NC) (QIAGEN, China), and siRNA-PUM2 (Genomeditech, China) were transfected to cells using the Lipofectamine™ 3000 Transfection Reagent according to the manufacturer's instructions (Invitrogen, USA).

MiRNA Pull-down and RNA Immunoprecipitation Chip (RIP) assays

The miRNA Pull-down assay was performed as previously described.<sup>38,39</sup> The RNA Immunoprecipitation Chip (RIP) assay was carried out using the Magna RIP RNA-Binding Protein Immunoprecipitation Kit according to the manufacturer's instructions (Millipore SiGMA, USA).

## Xenograft Mouse Model

Male C57BL/6 athymic nude mice (6 weeks old; Jackson Labs, USA) were housed in a pathogen-free animal room at  $24^\circ\text{C}$  with a 12 h dark-light cycle in the medicine animal facilities of Linyi chest Hospital under ad libitum feeding conditions. Mice were subcutaneously injected with a mixture of A549 cells ( $2 \times 10^5$ ) with PUM2-expressing or PUM2-knockdown CAFs ( $1 \times 10^5$ ) into the left and right buttocks ( $n=5$ ). Tumor volume was monitored every three days. Tumor volume was calculated using the formula:  $V = (L \times W^2) \times 0.5$ . After 24 days, mice were sacrificed and the tumor weight was measured.

## Statistical Analysis

Data were shown as means  $\pm$  SEM. The statistical analyses were performed using SPSS 19.0 software (SPSS, USA) and GraphPad Prism (GraphPad Software, USA). The differences between two or more groups were analyzed using one-way ANOVA or Student's *t*-test. Tukey's HSD test was performed for pairwise comparisons when the interaction was significant.  $P < 0.05$  was considered statistically significant.

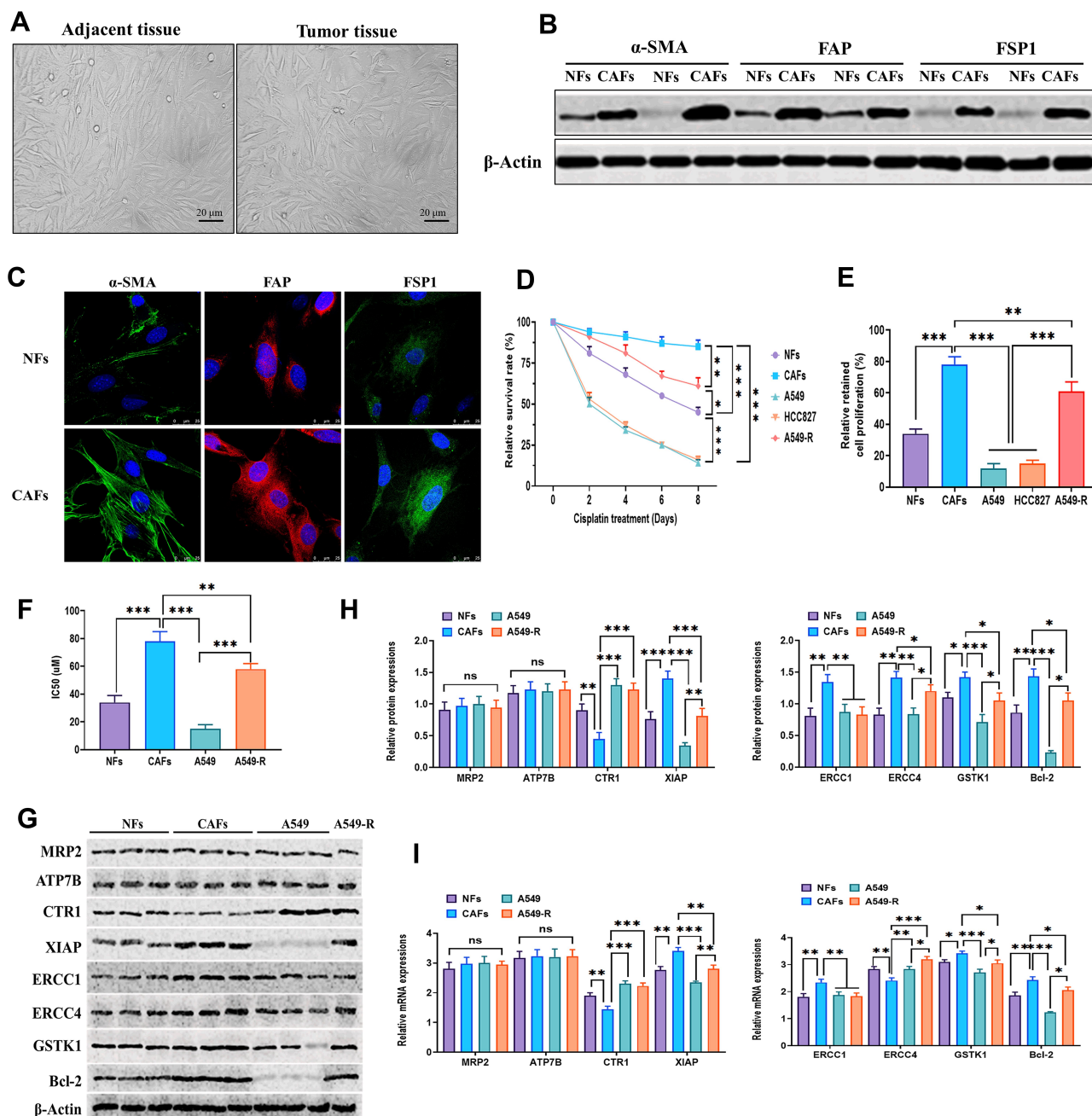
## Results

### NSCLC-Derived CAFs are Chemoresistant to Cisplatin

The CAFs from NSCLC tissues and their adjacent tissues were first isolated. The morphology of CAFs and

NFs displayed a spindle-like shape and were adherent in culture (Figure 1A). Western blotting immunofluorescence staining assays revealed that CAFs expressed higher levels of the fibroblast markers,  $\alpha$ -SMA, FAP, and FSP1, relative to NFs (Figure 1B and C). After treating with the same dosage of cisplatin, CAFs had

a higher survival rate than those of NFs, A549, HCC827, and cisplatin-resistant A549 cells (A549-R) (Figure 1D). Also, CAFs retained the highest level of proliferation in response to cisplatin treatment, compared with those of NFs and A549-R, while the most cisplatin-vulnerable cell lines, A549 and HCC827,

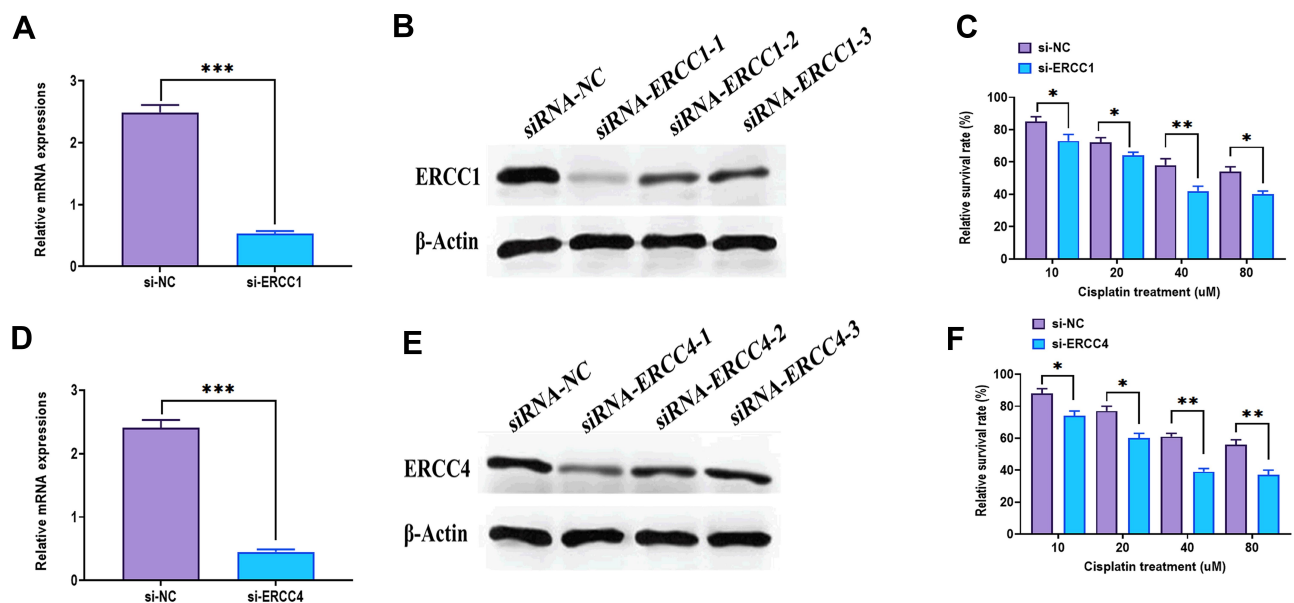


**Figure 1** NSCLC-derived CAFs are chemoresistant to cisplatin. (A) The morphology of NFs and CAFs under inverted microscopy. Scale bar: 20  $\mu$ m. (B) Western blots for specific fibroblast markers  $\alpha$ -SMA, FAP, and FSP1 in NFs and CAFs. Scale bar: 25  $\mu$ m. (C) Immunofluorescence assay for specific fibroblast markers  $\alpha$ -SMA, FAP, and FSP1 in NFs and CAFs. Scale bar: 25  $\mu$ m. (D) NFs, CAFs, A549, HCC827, and A549-R cells were treated with or without cisplatin (10  $\mu$ M) for 7 days, and cell viability was determined using the MTT assay. The relative survival rate was calculated as the ratio of obtained percent cell survival to that of control cells. (E) NFs, CAFs, A549, HCC827, and A549-R cells were treated with or without cisplatin (10  $\mu$ M) for 24 hours, and cell viability was determined by the percentage of proliferating cells after cisplatin treatment. (F) IC50 of cisplatin in NFs, CAFs, A549, and A549-R cells. (G and H) Protein expressions of cisplatin resistance-associated factors in NFs, CAFs, A549, and A549-R cells. (I) mRNA expressions of cisplatin resistance-associated factors in NFs, CAFs, A549, and A549-R cells. \* $P$  < 0.05; \*\* $P$  < 0.01; \*\*\* $P$  < 0.001.

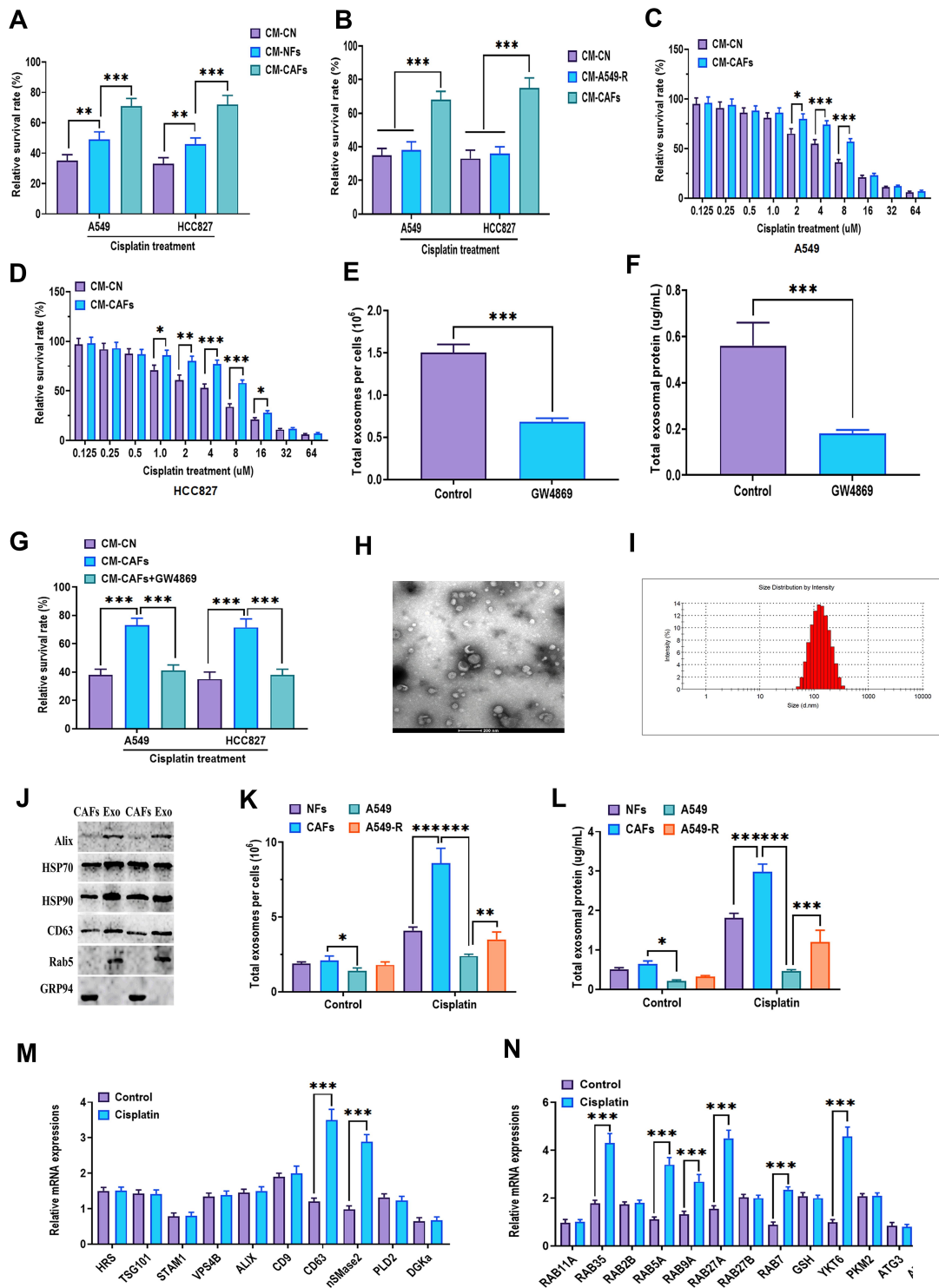
showed the lowest retained cell proliferation (Figure 1E). Next, the highest cisplatin IC<sub>50</sub> was found in CAFs, and A549 cells had the lowest IC<sub>50</sub> value (Figure 1F). The mechanism underlying cisplatin resistance is associated with multiple processes, such as reduced drug uptake, increased drug efflux or detoxification, alteration of DNA repair, or antiapoptotic proteins.<sup>40,41</sup> Next, Western blots were used to detect the protein expressions of several cisplatin resistance-associated regulators. As shown in Figure 1G and H, the protein expressions of XIAP, ERCC1, ERCC4, GSTK1, and Bcl-2 were higher in CAFs than those of NFs, A549, and A549-R cells, whereas CAFs had the lowest protein level of CTR1 than the other three cell types. Also, the similar mRNA expression profile of these genes was observed in CAFs, NFs, A549, and A549-R cells (Figure 1I). The ERCC1/4 system, nucleotide excision repair factors, displays a negative correlation with cisplatin-based survival response in NSCLC cells.<sup>42,43</sup> In this study, ERCC1 and ERCC4 were knockdown by specific siRNAs (Figure 2A, B, D, and E), and the knockdown of ERCC1 or ERCC4 resulted in reduced cisplatin resistance in CAFs (Figure 2C and F). Collectively, these observations suggest that NSCLC-derived CAFs are innately chemoresistant to cisplatin.

## CAFs-Derived Exosomes Confer Cisplatin Resistance of NSCLC Cells

Given the innate cisplatin resistance of CAFs, further investigation of whether CAFs-derived bioactive factors could impact the cellular activities of NSCLC cells was conducted. The present study demonstrated that CAFs-conditioned medium (CM-CAFs) significantly increased the survival rate in A549 and HCC827 cells, compared with CM-NFs and CM-NSCLC cells (CM-CN) (Figure 3A). Also, A549-R-conditioned medium (CM-A549-R) did not exert a similar function to promote the survival rate in NSCLC cells in response to cisplatin treatment (Figure 3B). The effect of CM-CAFs was detected in NSCLC cells in a wide dosage range of cisplatin, namely, 2–8  $\mu$ M in A549 cells and 1–16  $\mu$ M in HCC827 cells (Figure 3C and D). Exosomes have emerged as essential message-transporters between CAFs and cancer cells.<sup>27,44</sup> Thus, it was speculated that CAFs-derived exosomes might play an essential role in the effect of CM-CAFs on NSCLC cells. First, GW4869, a neutral sphingomyelinase inhibitor for blocking exosome generation<sup>45</sup> was applied to CM-CAFs, which showed that GW4869 significantly reduced the number of exosomes as well as exosomal protein in CM-CAFs (Figure 3E and F). Furthermore, the results showed that the addition of GW4869 abolished the effect CM-CAFs on both A549 and



**Figure 2** Expressions of ERCC1 and ERCC4 in CAFs. (A) mRNA expression of ERCC1 in CAFs transfected with siRNA-ERCC1. (B) Protein expression of ERCC1 in CAFs transfected with siRNA-ERCC1. (C) Cell viability of CAFs with ERCC1 knockdown in response to different concentrations of cisplatin. (D) mRNA expression of ERCC1 in CAFs transfected with siRNA-ERCC1. (E) Protein expression of ERCC1 in CAFs transfected with siRNA-ERCC1. (F) Cell viability of CAFs with ERCC1 knockdown in response to different concentrations of cisplatin. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .



**Figure 3** CAFs-derived exosomes confer cisplatin resistance of NSCLC cells. **(A)** Cell viability of A549 and HCC827 cells cultured in CM-NFs or CM-CAFs after cisplatin treatment. Control mediums (CM-CN) were A549 or HCC827 cells' own culture mediums. Cell viability was determined by MTT assays. The relative survival rate was calculated as the ratio of obtained percent cell survival to that of control cells. **(B)** Cell viability of A549 and HCC827 cells cultured in CM-A549-R or CM-CAFs after cisplatin treatment. Control mediums (CM-CN) were A549 or HCC827 cells' own culture mediums. **(C and D)** Cell viability of A549 and HCC827 cells cultured in CM-CAFs in response to different concentrations of cisplatin. Control mediums (CM-CN) were A549 or HCC827 cells' own culture mediums. **(E)** The number of exosomes in CM-CAFs treated with 20  $\mu$ M GW4869 after cisplatin treatment. Control mediums (CM-CN) were A549 or HCC827 cells' own culture mediums. **(F)** Exosomal protein concentration in CM-CAFs treated with 20  $\mu$ M GW4869. **(G)** Cell viability of A549 and HCC827 cells cultured in CM-CAFs or CM-CAFs plus 20  $\mu$ M GW4869 after cisplatin treatment. Control mediums (CM-CN) were A549 or HCC827 cells' own culture mediums. **(H)** The morphology of CAFs-derived exosomes, as imaged by transmission electron microscopy. **(I)** The size distributions and the number of CAFs-derived exosomes, as determined by nanoparticle tracking analysis. **(J)** Protein expressions of exosome markers Alix, HSP90, HSP70, CD63, and Rab5 and negative marker cellular proteins, GRP94. **(K)** The number of exosomes in NFs, CAFs, A549, and A549-R cells treated with or without cisplatin treatment. **(L)** Exosomal protein concentration in NFs, CAFs, A549, and A549-R cells treated with or without cisplatin treatment. **(M and N)** mRNA expressions of exosome generation-associated factors. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .



HCC827 cells under cisplatin treatment (Figure 3G), which suggests that exosomes participate in CM-CAFs-induced cisplatin resistance in NSCLC cells. Next, the exosomes were isolated from CM-CAFs through the ultracentrifugation assay and nanoparticle tracking analysis. Transmission electron microscopy was applied to determine the morphology and size of the exosomes. As shown in Figure 3H and I, CAFs-derived exosomes displayed typical spherical shape and were 90–130 nm in diameter. Also, Western blots revealed that CAFs-derived exosomes positively expressed exosome markers, including Alix, HSP70, HSP90, CD63, and Rab5, but not the negative marker, GRP94 (Figure 3J). Notably, compared with NFs, A549, and A549-R, CAFs generated the highest number of exosome and exosomal protein in the presence of cisplatin treatment (Figure 3K and L). Given the complexity of the biogenesis of exosomes, this multi-step process includes multivesicular bodies (MVBs) transportation, docking, and fusion with plasma membranes, of which several essential molecule families, such as SNARE and RAB, play an essential role.<sup>46,47</sup> In the present study, mRNA expressions of several critical regulators for the generation of exosomes were determined and the results showed that the mRNA expressions of CD63, nSMase2, RAB35, RAB5A, RAB9A, RAB27A, RAB7, and YKT6 were increased in cisplatin-treated CAFs (Figure 3M and N), further indicating that cisplatin may activate an exosome-generation system in CAFs in response to cisplatin treatment.

## CAFs-Derived Exosomes are Taken Up by NSCLC Cells

To determine whether CAFs-derived exosomes were taken up by A549 and HCC827 cells, exosomes were labeled with DiO and cocultured with A549 and HCC827 cells for 24 hours. As shown in Figure 4A, DiO-labeled green exosomes were taken up by both A549 and HCC827 cells. By applying the same protocols mentioned above, the characteristics of exosomes derived from CAFs control, cisplatin-treated CAFs and NFs were verified, respectively (Figure 4B). Subsequently, the results revealed that exosomes derived from cisplatin-treated CAFs displayed a more potent effect on the survival rate of A549 and HCC827 cells in the presence of cisplatin, compared with exosomes derived from CAFs without cisplatin treatment (Figure 4C) and A549-R cells (Figure 4D) and NFs with cisplatin treatment (Figure 4E). Together, these data suggest that CAFs or cisplatin-

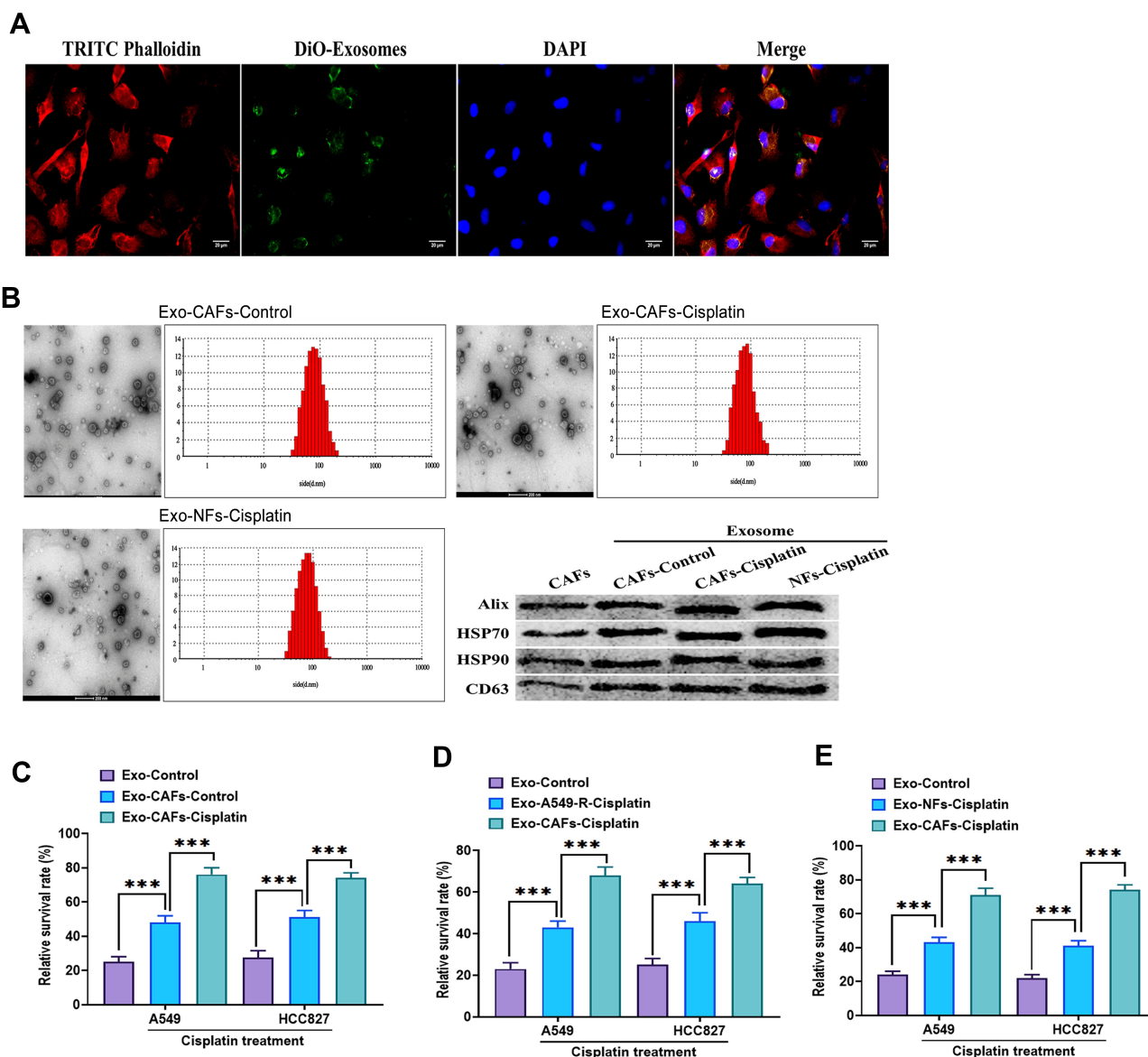
treated CAFs-derived exosomes were internalized by NSCLC cells and promoted the survival rate after cisplatin treatment.

## Exosomal miRNA-130a is Transferred from CAFs to NSCLC Cells

Accumulating evidence suggests that exosomes can transfer a group of bioactive molecules such as miRNAs, lncRNAs, and proteins from CAFs to cancer cells, influencing cellular activities of recipient cells.<sup>23,27</sup> Next, miRNA array analysis was used to determine the functional molecules, particularly miRNAs that contributed to the effect of CAFs-derived exosomes on NSCLC cells. The results showed that miRNA-130a significantly increased CAFs-derived exosomes, compared with those derived from CAFs without cisplatin treatment (Figure 5A). Meanwhile, real-time PCR revealed that the expression of miRNA-130a was higher in cisplatin-treated CAFs than in CAFs without cisplatin treatment (Figure 5B). Also, the level of miRNA-130a was higher in A549 and HCC827 cells treated with exosomes derived from cisplatin-treated CAFs, relative to those treated with exosomes derived from NSCLC cells own mediums (Figure 5C). Intriguingly, miRNA-130a was increased in NSCLC tissue relative to adjacent normal tissues (Figure 5D). Furthermore, CAFs innately expressed a higher level of miRNA-130a compared with those of NFs, A549, HCC827, and A549-R cells (Figure 5E).

To investigate whether miRNA-130a was transferred from CAFs to NSCLC cells through exosomes, A549 and HCC827 cells were cultured in CM-CAFs or CM-CAFs with the addition of GW4869. The results showed that the expression of miRNA-130a was increased in A549 and HCC827 cells cultured in CM-CAFs, while the increased level of miRNA-130a was reversed by the addition of GW4869 (Figure 5F). Additionally, CAFs transfected with Cy3-labeled miRNA-130a were cocultured with A549 and HCC827 cells for 48 hours and Cy3-labeled green miRNA-130a was found in NSCLC cells (Figure 5G). Also, the expression of miRNA-130a did not change in CM-CAFs with RNaseA, while was dramatically decreased when Triton X-100 was added to CM-CAFs (Figure 5H), suggesting that miRNA-130a was primarily encased within the membrane structures. Moreover, there was no difference in the level of miRNA-130a between CM-CAFs and CAFs-derived exosomes (Figure 5I). Collectively, miRNA-130a was transferred from CAFs to recipient NSCLC cells through exosomes. To determine

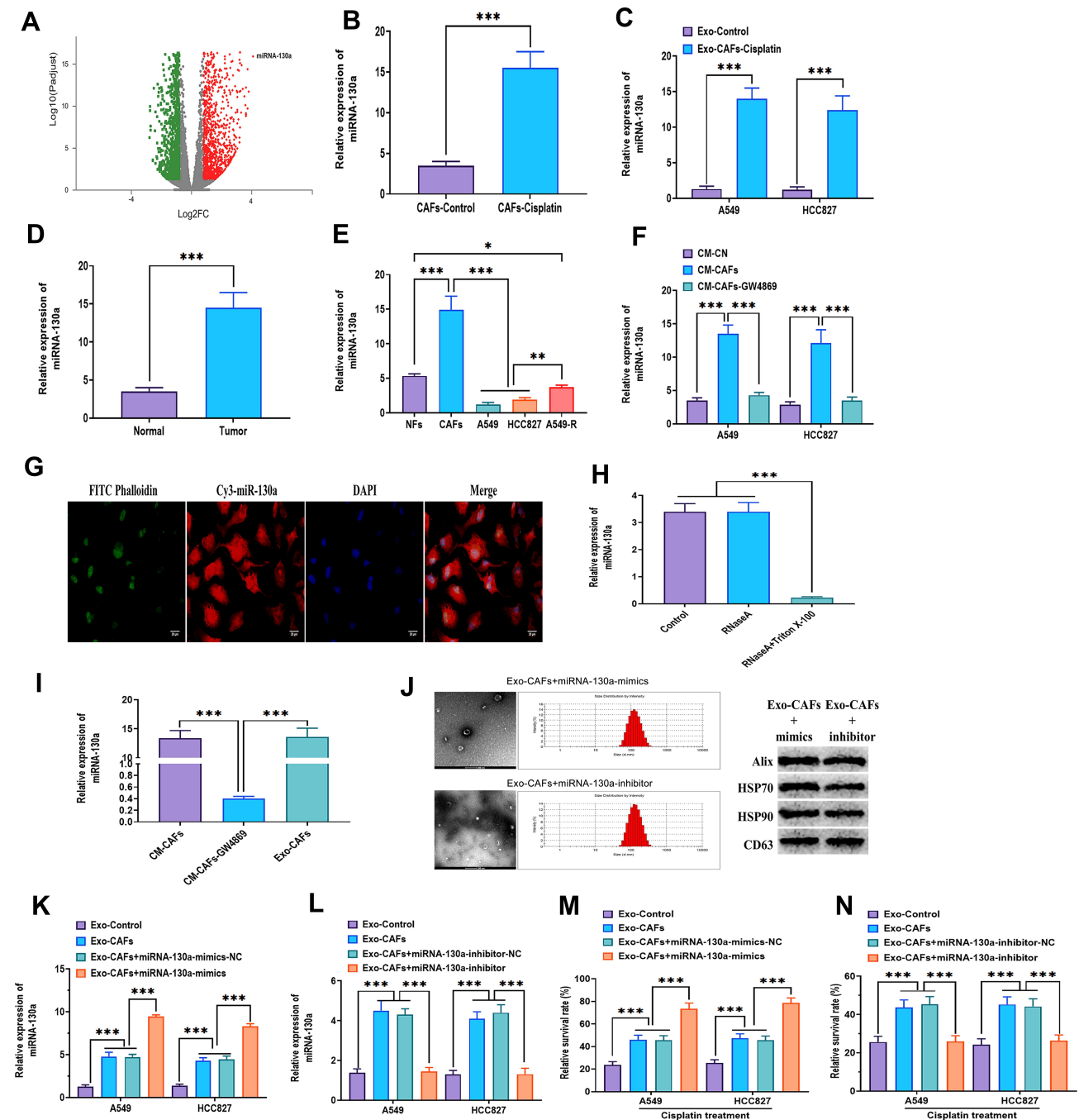




**Figure 4** CAFs-derived exosomes were taken up by NSCLC cells. **(A)** A549 cells were incubated with CAFs-derived exosomes labeled with DiO (25  $\mu\text{g}/\text{mL}$ ) for 24 hours, and the image was taken under confocal microscopy. TRITC Phalloidin (red) indicated cytoskeleton. DiO (green) indicated exosome. DAPI (blue) indicated the nuclei. Scale bar: 20  $\mu\text{m}$ . **(B)** The characteristics of exosomes were identified by transmission electron microscopy, nanoparticle tracking analysis, and Western blots. **(C)** Cell viability of A549 and HCC827 cells treated with exosomes (25  $\mu\text{g}/\text{mL}$ ) derived from CAFs or cisplatin-treated CAFs. Control exosomes (Exo-control) derived from A549 or HCC827 cells' own culture mediums. Cell viability was determined by MTT assays. The relative survival rate was calculated as the ratio of obtained percent cell survival to that of control cells. **(D)** Cell viability of A549 and HCC827 cells treated with exosomes (25  $\mu\text{g}/\text{mL}$ ) derived from cisplatin-treated A549-R cells or CAFs. Control exosomes (Exo-control) derived from A549 or HCC827 cells' own culture mediums. **(E)** Cell viability of A549 and HCC827 cells treated with exosomes (25  $\mu\text{g}/\text{mL}$ ) derived from cisplatin-treated NFs or CAFs. Control exosomes (Exo-control) derived from A549 or HCC827 cells' own culture mediums. \*\*\* $P < 0.001$ .

the function of miRNA-130a in CAFs-derived exosomes, CAFs were transfected with miRNA-130a mimics and inhibitors and the exosomes were isolated and then identified using the same methods mentioned above (Figure 5J). The expression level of miRNA-130a was increased and decreased in exosomes derived from CAFs with overexpression of miRNA-130a or miRNA-130a knockdown, respectively (Figure 5K and L). Functionally, the survival

rate was increased in NSCLC cells treated with exosomes derived from CAFs or CAFs with overexpression of miRNA-130a, while exosomes derived from CAFs with miRNA-130a knockdown abolished this effect (Figure 5M and N). These results collectively suggest that CAFs-derived exosomes promote the survival rate of NSCLC cells after cisplatin treatment through exosomal miRNA-130a.



**Figure 5** Exosomal miRNA-130a is transferred from CAFs to NSCLC cells. **(A)** Volcano Plotting of differentially expressed miRNAs between NFs- and CAFs-derived exosomes. miRNAs had  $\geq 2$ -fold change and adjusted  $P < 0.05$  were considered as differentially expressed miRNAs. **(B)** Expression of miRNA-130a in CAFs and cisplatin-treated CAFs. **(C)** Expression of miRNA-130a in A549 and HCC827 cells treated with exosomes derived from cisplatin-treated CAFs (25  $\mu\text{g}/\text{mL}$ ). Control exosomes (Exo-control) derived from A549 or HCC827 cells' own culture mediums. **(D)** Expression of miRNA-130a in normal and NSCLC tumor tissues. **(E)** Expression of miRNA-130a in NFs, CAFs, A549, HCC827, and A549-R cells. **(F)** Expression of miRNA-130a in A549 and HCC827 cells cultured in CM-CAFs or CM-CAFs plus 20  $\mu\text{M}$  GW4869. Control mediums (CM-CN) were A549 or HCC827 cells' own culture mediums. **(G)** A549 cells were incubated with CAFs transfected with Cy3-tagged miRNA-130a for 48 hours, and the image was taken under confocal microscopy. FITC Phalloidin (green) indicated cytoskeleton. Cy3 (red) indicated exosome. DAPI (blue) indicated the nuclei. Scale bar: 20  $\mu\text{m}$ . **(H)** Expression of miRNA-130a in CM-CAFs treated with RNaseA or RNaseA plus Triton X-100. **(I)** Expression of miRNA-130a in CM-CAFs, CM-CAFs plus 20  $\mu\text{M}$  GW4869, or CAFs-derived exosomes. **(J)** The characteristics of exosomes were identified by transmission electron microscopy, nanoparticle tracking analysis, and Western blots. **(K)** Expression of miRNA-130a in exosomes derived from CAFs or CAFs treated with miRNA-130a mimics. **(L)** Expression of miRNA-130a in exosomes derived from CAFs or CAFs treated with miRNA-130a inhibitors after cisplatin treatment. Control exosomes (Exo-control) derived from A549 or HCC827 cells' own culture mediums. **(M)** Cell viability of A549 and HCC827 cells treated with exosomes derived from CAFs or CAFs treated with miRNA-130a mimics after cisplatin treatment. Control exosomes (Exo-control) derived from A549 or HCC827 cells' own culture mediums. Cell viability was determined by MTT assays. The relative survival rate was calculated as the ratio of obtained percent cell survival to that of control cells. **(N)** Cell viability of A549 and HCC827 cells treated with exosomes derived from CAFs or CAFs treated with miRNA-130a inhibitors after cisplatin treatment. Control exosomes (Exo-control) derived from A549 or HCC827 cells' own culture mediums. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .

## Exosomal miRNA-130a Packaging is Associated with PUM2

To determine the mechanism underlying the packaging of exosomal miRNA-130a in CAFs, a database of RBP specificities (RBPDB)<sup>48</sup> was used to analyze the interaction between miRNA-130a and motifs of RNA-binding proteins (RBPs). The results revealed that PUM2 and eukaryotic translation initiation factor 4B (EIF4B) motifs displayed specific miRNA-130a binding sites (Figure 6A). Next, specific siRNAs were applied to knockdown PUM2 and EIF4B in CAFs (Figure 6B and C) and showed that PUM2 knockdown decreased the level of exosomal miRNA-130a while the level of cellular miRNA-130a did not change (Figure 6D and F). Also, knockdown of EIF4B did not alter the levels of exosomal miRNA-130a (Figure 6E). Furthermore, the mechanical interactions between miRNA-130a and PUM2 in both cytoplasm and exosomes, but not in the nucleus, and these interactions were disrupted when the sequence of miRNA-130a was mutated (Figure 6G). Moreover, the RIP assay was performed in cell and exosome lysates of CAFs which showed that miRNA-130a enriched the PUM2 antibody, relative to the IgG group (Figure 6H). Also, there was more miRNA-130a enrichment in cell and exosome lysates of cisplatin-treated CAFs compared with those of CAFs without cisplatin treatment (Figure 6I). Functionally, the level of exosomal miRNA-130a was decreased in CAFs with PUM2 knockdown (Figure 6J). In addition, cisplatin increased the level of PUM2 protein in exosomes derived from CAFs but did not alter PUM2 protein expression in CAFs (Figure 6K). Collectively, these findings indicate that cisplatin treatment may promote the packaging of miRNA-130a in exosomes of CAFs through the mediation of PUM2.

## Effect of CAFs-Derived PUM2 on Tumor Growth in vivo

To determine the effect of CAFs-derived PUM2 on tumor growth, A549 plus CAFs with PUM2 overexpression/knockdown was used in xenograft mice (Figure 7A). The results showed that A549 plus CAFs with PUM2 overexpression was associated with larger tumor weights and faster tumor growth (Figure 7B–D), whereas tumors formed by A549 plus CAFs with PUM2 knockdown displayed smaller tumor weights and slower growth (Figure 7G–I). Also, Western blots revealed that the level of PUM2 was upregulated in tumor tissues treated with A549 plus CAFs with PUM2 overexpression (Figure 7E and F), while was

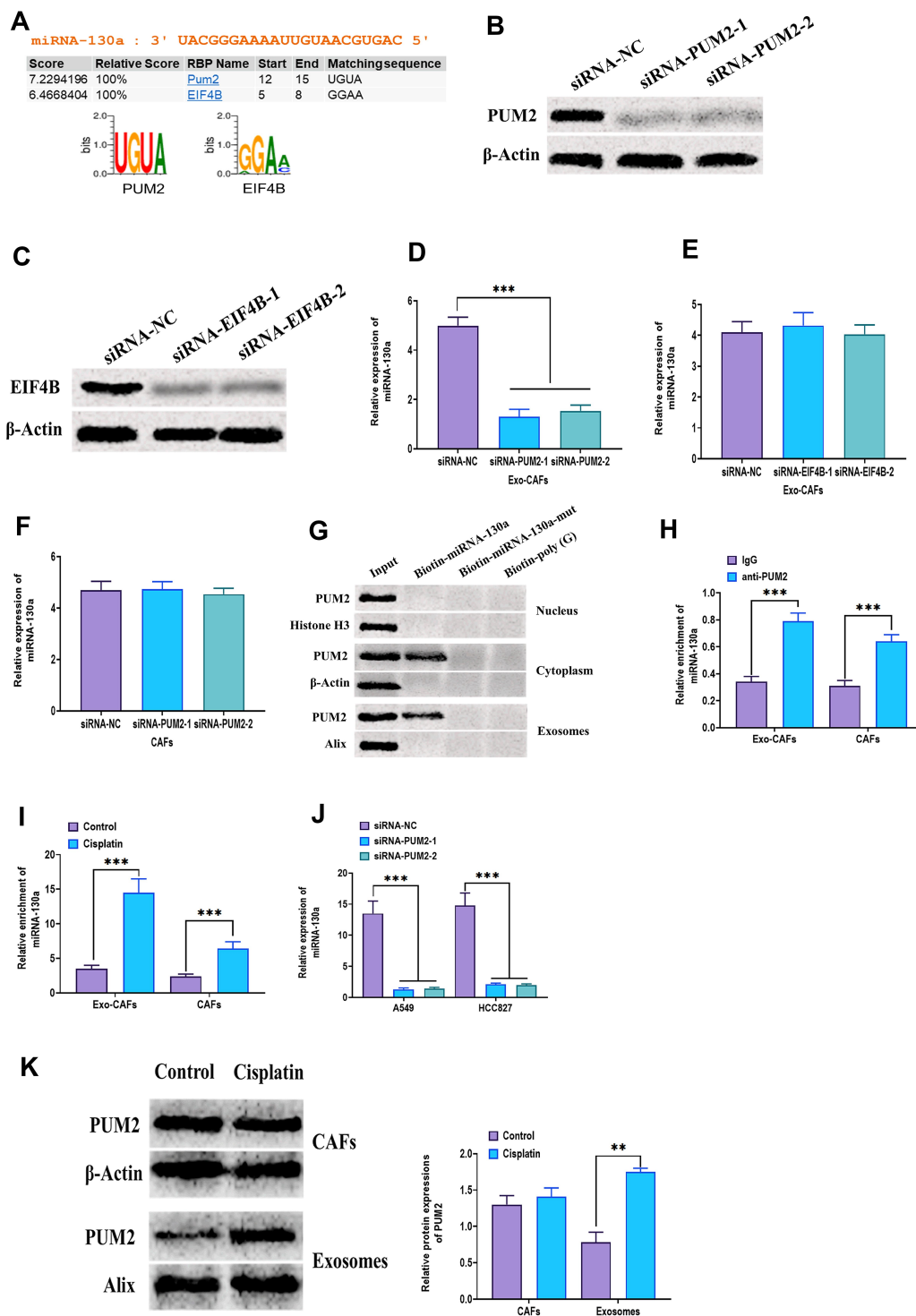
downregulated in those treated with A549 plus CAFs with PUM2 knockdown (Figure 7J and K).

## Discussion

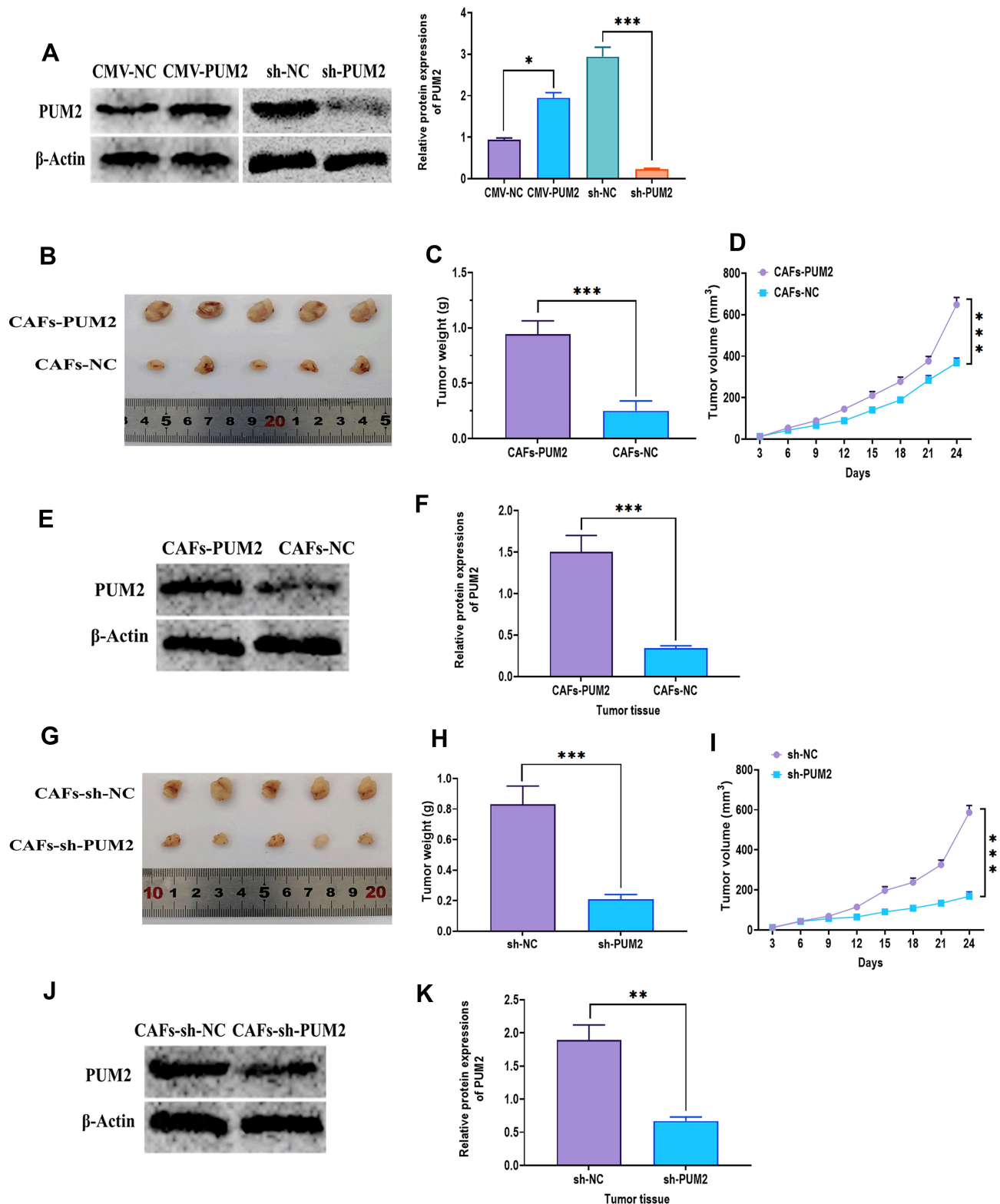
Chemoresistance is one of the primary challenges for the treatment and management of cancers. Given its heterogeneous nature, the mechanism related to chemoresistance is complex and is not comprehensively understood, thereby cancer cell resistance to chemotherapeutic agents continues to pose a significant barrier for all oncologists and their patients.<sup>49</sup> A previous study revealed that nearly 63% (1409/2227) and 68% (1056/1565) of NSCLC patients' tumor cultures are extreme or intermediate resistant to cisplatin and carboplatin, respectively.<sup>50</sup> Beyond that, these NSCLC samples display a different degree of resistance to other drugs, including doxorubicin, gemcitabine, docetaxel, vinorelbine, etoposide, paclitaxel, and topotecan.<sup>50</sup> Thus, all NSCLC patients will ultimately develop chemoresistance to drugs, even with a satisfactory initial response.<sup>51</sup> To overcome this barrier, there is an urgent need to develop novel chemotherapy agents as well as to explore the molecular mechanism behind the chemoresistance of NSCLC.

Accumulating evidence has demonstrated that CAFs are the most abundant stromal cells in the tumor microenvironment, playing an essential role in the response of cancer cells to chemotherapy.<sup>52,53</sup> In the present study, NSCLC-derived CAFs are innately resistant to cisplatin treatment and that CAFs-CM significantly improved the survival rate and proliferation of NSCLC cell lines, A549 and HCC827, in the presence of cisplatin. Furthermore, after coculturing with CAFs, A549 and HCC827 cells were insensitive to cisplatin. These observations together suggest that CAFs confers with NSCLC cells on the resistance to chemotherapeutic agents. A similar role of CAFs in chemoresistance has also been reported in other cancer types, such as ovarian cancer,<sup>54</sup> colorectal cancer,<sup>55</sup> and gastric cancer.<sup>56</sup> However, due to the complexity and heterogeneity of intercellular communication between CAFs and cancer cells, the molecular mechanism underlying how CAFs regulate chemotherapy resistance in cancers is still not well documented.

Growing evidence has suggested that CAFs can impact cancer progression through various bioactive factors, including growth factors,<sup>57</sup> cytokines,<sup>31</sup> and chemokines.<sup>58</sup> In addition, exosome-mediated intercellular interaction between CAFs and cancer cells has been reported to be involved in the chemoresistance of tumor cells.<sup>59,60</sup> In the present study, we found that CAFs-derived exosomes promoted the survival rate of NSCLC cells in response to cisplatin and that exosome



**Figure 6** Exosomal miRNA-130a packaging is associated with PUM2. **(A)** Prediction of interaction between miRNA-130a and RBP motifs. **(B)** Protein expression of PUM2 in CAFs treated with siRNA-PUM2. **(C)** Protein expression of EIF4B in CAFs treated with siRNA-EIF4B. **(D)** Expression of miRNA-130a in exosomes derived from CAFs treated with siRNA-PUM2. **(E)** Expression of miRNA-130a in exosomes derived from CAFs treated with siRNA-EIF4B. **(F)** Expression of miRNA-130a in CAFs treated with siRNA-PUM2. **(G)** Protein expression of PUM2 in samples from miRNA pull-downs assay. MiRNA pull-downs assay was carried out with CAFs nuclear, cytoplasmic, or exosomal lysates and the biotinylated miRNA-130a or mutated miRNA-130a. **(H)** RNA Immunoprecipitation Chip (RIP) assay was performed with anti-PUM2 antibody and IgG on CAFs or CAFs-derived exosomes. The expression of miRNA-130a was detected by real-time PCR in immunoprecipitated samples. **(I)** RIP assay was performed to determine the level of miRNA-130a in CAFs cytoplasmic or exosomal lysates with or without cisplatin treatment. **(J)** Expression of miRNA-130a in A549, HCC827 cells treated with siRNA-PUM2. **(K)** Protein expression of PUM2 in CAFs or CAFs-derived exosomes with or without cisplatin treatment. \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .



**Figure 7** Effect of CAFs-derived PUM2 on tumor growth in vivo. **(A)** Protein expression of PUM2 in CAFs cells transfected with PUM2-expressing or PUM2-silencing plasmid for 48 hours. **(B)** Representative image of tumor tissues of nude mice subcutaneously injected with a mixture of A549 cells plus CAFs transfected with PUM2-expressing plasmid. **(C and D)** Tumor weight and growth rate of nude mice subcutaneously injected with a mixture of A549 cells plus CAFs transfected with PUM2-expressing plasmid. **(E and F)** Protein expression of PUM2 in tumor tissues of nude mice subcutaneously injected with a mixture of A549 cells plus CAFs transfected with PUM2-expressing plasmid. **(G)** Representative image of tumor tissues of nude mice subcutaneously injected with a mixture of A549 cells plus CAFs transfected with PUM2-silencing plasmid. **(H and I)** Tumor weight and growth rate of nude mice subcutaneously injected with a mixture of A549 cells plus CAFs transfected with PUM2-silencing plasmid. **(J and K)** Protein expression of PUM2 in tumor tissues of nude mice subcutaneously injected with a mixture of A549 cells plus CAFs transfected with PUM2-silencing plasmid. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .



secretion inhibitor, GW4869, significantly reversed the effect of CAFs-derived exosomes on chemoresistance of NSCLC cells. Numerous studies reported that CAFs-derived exosomes are essential for the regulation of chemoresistance in various cancers, including bladder cancer,<sup>61</sup> ovarian cancer,<sup>62</sup> and colorectal cancer.<sup>55</sup> Furthermore, cisplatin treatment significantly upregulated the generation of exosomes from CAFs. Therefore, these findings reported may provide a novel CAFs-targeting therapeutic strategy for NSCLC.

To further investigate the mechanism by which CAFs-derived exosomes promoted cisplatin resistance of NSCLC cells, miRNA array analysis was performed to determine the functional miRNAs that contributed to the effect of CAFs-derived exosomes on NSCLC cells. In the present study, CAFs-derived exosomal miRNA-130a played an important role in endowing cisplatin resistance to NSCLC. As a multi-functional factor, miRNA-130a participates in various cellular activities in cancer cells, for example, autophagy,<sup>63</sup> proliferation,<sup>64</sup> and apoptosis.<sup>65</sup> MiRNA-130a has been reported to be an oncogenic factor, whose upregulation in NSCLC tissue is positively correlated to poor prognosis, lymph node metastasis, and stage of tumor node metastasis.<sup>66</sup> Meanwhile, miRNA-130a has been reported to play an important role in chemoresistance of various cancer types.<sup>67</sup> Particularly, miRNA-130a is upregulated in hepatoma cells in the presence of cisplatin, which is associated with the involvement of RUNX3/Wnt signaling.<sup>68</sup> Also, the downregulation of miRNA-130a can attenuate the cisplatin resistance in ovarian cancer cells through MDR1/P-glycoprotein pathways.<sup>69</sup> In another study reported by Zhang et al, X-linked inhibitor of apoptosis (XIAP) is identified to be a direct targeting gene of miRNA-130a to confer cisplatin resistance in ovarian cancer.<sup>70</sup> These findings collectively demonstrate that the role of miRNA-130a may be involved in a complex regulatory network and displays a cancer type-dependent manner. Given its important role in chemoresistance, miRNA-130a is a potential therapeutic target for developing new drugs for NSCLC treatment.

Given the upregulation of miRNA-130a in CAFs-derived exosomes after cisplatin treatment, the next goal was to investigate how miRNA-130a was packaged into exosomes. It has been demonstrated that extracellular miRNAs are encased in small-size membranous vesicles or packaged with RNA-binding proteins.<sup>71</sup> Through bioinformatics analysis, two RBPs, PUM2 and EIF4B, had binding sequences to bind miRNA-130a. Further functional experiments demonstrated that PUM2 knockdown, not EIF4B, in CAFs

significantly downregulated the level of miRNA-130a in CAFs-derived exosomes and recipient NSCLC cells after cisplatin treatment. Meanwhile, the upregulation of PUM2 promoted tumor growth of xenograft mice while knockdown of PUM2 exerted the opposite role. These results suggested that PUM2 may play an essential role in packaging miRNA-130a into exosomes, thereby indirectly impacting the abundance and function of miRNA-130a in CAFs, exosomes, and NSCLC cells. Thus, PUM2 may act as a target to inhibit the oncogenic effect of miRNA-130.

## Conclusion

In conclusion, the results demonstrate that NSCLC-associated CAFs were innately resistant to cisplatin, and CAFs-derived exosomes conferred chemoresistance of NSCLC cells through transferring miRNA-130a. Also, PUM2 was associated with packaging miRNA-130a into exosomes. Thus, CAFs-derived exosomal miRNA-130a may be a potential therapeutic target for chemoresistance in NSCLC.

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## Disclosure

The authors declare no conflicts of interest.

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