ORIGINAL RESEARCH

A Super-Assembled Synergistically Nanoplatform AP@ZIF-8^{Pt} for Hepatocarcinoma Therapy

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Introduction: Intensive cancer treatment with nanoplatform is widely exploited in the clinic, the emerging nanomedicine offers an unparalleled opportunity for encapsulating potential antitumor drugs in a nano-carrier. Apoptin (AP), a coding protein of VP3 gene, stem from the chicken anemia virus (CAV), can be activated in malignant cells selectively and prevents the dividing cancer cells from repairing their DNA lesions, thereby forcing them to undergo apoptosis. Herein, a three-step intelligent biodegradable drug delivery nanoplatform was designed.

Methods: First, a hollow ZIF-8 was synthesized, embedded with platinum nanoparticle to form $ZIF-8^{Pt}$, and then loaded with AP, and lastly formed AP@ZIF-8^{Pt}, which possess pH-responsive drug release and cancer-targeted ability.

Results: As expected, both in vitro and in vivo experiment demonstrated that AP@ZIF-8^{Pt} performed treatment effects in hepatocarcinoma through relieving tumor-hypoxic microenvironment, inhibiting cell proliferation and promoting cell apoptosis. Further transcriptomic analysis showed that the specific mechanism of the AP@ZIF-8^{Pt} was thermogenesis, signaling pathways regulating pluripotency of stem cells, ribosome, prion disease and PI3K-Akt signaling pathway.

Discussion: This work highlights a new strategy for liver cancer treatment and provides a reference for treating malignant tumors. **Keywords:** AP@ZIF-8^{Pt}, tumor-hypoxic microenvironment, apoptosis, PI3K-Akt signaling pathway

Introduction

Liver cancer, specifically hepatocellular carcinoma (HCC), is one of the most frequent malignancies that causes cancerrelated death.¹⁻³ Although immune checkpoint blockade-based immunotherapies and targeted therapy of HCC have achieved remarkable success in the field of HCC treatment, few of them induce durable responses and provide prominent survival benefits in patients because of the HCC physiological complexity.⁴⁻⁶ To overcome it, effective strategies to optimizing HCC treatment are urgently needed,^{7,8} and there are two emerged strategies: finding new drugs according to specific mechanisms and developing novel drug delivery system to give full play to the new drugs.9

As for the new drugs finding, drug selectivity to cancer cells is the key issue of antitumor therapy.¹⁰ A group of proteins, killing cancer cells specifically without harming the normal cells, has attracted scientific interest.¹¹ Apoptin (AP) is one of these proteins. In the normal cells, AP becomes filamentous to aggregated and then degraded through proteasomes. While in cancer cells, AP induces apoptosis.¹²⁻¹⁴ Our early study has proved that AP could induce apoptosis in hepatocarcinoma cells through targeting XPO1.¹⁵ So we focused on AP as the potential HCC therapy. As for the novel drug delivery system developing, the emerging nanoplatform provides a new direction, which delivering the drugs and regulating tumor-hypoxic microenvironment synergistically, therefore achieving remarkable therapeutic effect.¹⁶ Among them, zeolitic imidazolate framework-8 (ZIF-8) is a metal-organic framework (MOF) serving as a promising candidate for its stability in aqueous environments and

Graphical Abstract



decomposability in acidic environments.^{17–19} Moreover, ZIF-8 possesses good biocompatibility, can reduce immune reactions and increase drugs bioavailability in vivo.^{20–22} Platinum nanoparticles (Pt) can catalyze H_2O_2 (100 M-1 mm) to produce oxygen, thereby relieving tumor hypoxia in tumor tissues.^{23,24}

Herein, we design a fire-new nanoplatform, using ZIF-8 implanted with $Pt(ZIF-8^{Pt})$, after AP loading, the final AP@ZIF-8^{Pt} was used to HCC therapy. First, AP@ZIF-8^{Pt} had decomposability in the weakly acidic tumor microenvironment (TME), and then releases loaded AP. Second, the released AP can induce apoptosis in hepatocarcinoma cells. Third, Pt works as a catalyst, trigger endogenous H₂O₂ into O₂, relieving hypoxia of TME and further improving the drug efficacy of AP. Expectedly, both in vitro and in vivo experiment results demonstrated that AP@ZIF-8^{Pt} possessed the extraordinary biocompatibility and enhanced therapeutic effect through inducing apoptosis, promoting DNA damage of hepatocarcinoma cells. Further transcriptomic analysis showed that the specific mechanism of the AP@ZIF-8^{Pt} was thermogenesis, signaling pathways regulating pluripotency of stem cells, ribosome, prion disease and PI3K-Akt signaling pathway. Collectively, this work highlights a new strategy for HCC therapy and can be a reference for developing the advanced antitumor therapy.

Materials and Methods

Materials

1,2-dimethylimidazole (2-MIL), zinc nitrate hexahydrate (Zn (NO₃)₂·6H₂O), chloroplatinic acid hydrate (H₂PtCl₆·6H₂O) and polyvinylpyrrolidone (PVP) were purchased from Aladdin. AP was synthesized by Shanghai Qiangyao Biotechnology Co., Ltd. 4'6-diamidino-2-phenylindole (DAPI) and 2,7-dichlorodihydrofluorescein diacetate (DCFH-DA) were purchased from Beyotime Biotechnology. The γ -H2AX antibody, HIF-1 α antibody and secondary antibodies were purchased from Cell Signaling Technology (Danvers, MA, USA). All of the reagents used in the experiments were of analytical grade and used directly without treatment.

Characterizations

Malvern Zetasizer Nano ZS90 Apparatus, UV-2550 Ultraviolet-visible Spectrophotometer (Shimadzu, Japan), transmission electron microscopy system (JEOL/JEM-F200), XRD (Bruker/D8 ADVANCE), FTIR (Bruker/VERTEX70), FC500 flow cytometer (Beckman Coulter, USA) were used for material characterization same as our previous studies.^{25,26}

Synthesis of AP@ZIF-8^{Pt}

Zinc nitrate hexahydrate (25 mm) and 2-methyl imidazole (200 mm) were first dissolved in methanol. Then, 20 mL of methanol solution was stirred together with Pt nanoparticles. After continuous stirring for 3–4 h, the sample was washed three times with methanol and ZIF-8^{Pt} was obtained. Then redispersed ZIF-8^{Pt} in ethanol (10 mL) added AP and stirred for 6–8 h. The mixed solution was then washed with methanol (10000 rpm, 10 min) to remove excess AP then obtained $AP@ZIF-8^{Pt}$.

In vitro Cytotoxicity Assay

The cytotoxicity against HCC cells was measured by CCK8 assay. The cells were cultured in a 96-well plate and incubated overnight. Next, gradient concentrations of AP@ZIF-8^{Pt} (0, 20, 40, 60, 80 and 100 μ g/mL) were added to each well and incubated for 24 h. Five parallel samples were set for each group, then 10 μ L of CCK8 was added to each well and incubated for 4 h. The absorbance was quantified by a spectrophotometer at 450 nm.

Reactive Oxygen Species (ROS) Detection

ROS in cells was detected using the ROS probe, DCFH-DA. Briefly, HCC cells were seeded in a 6-well plate and cultured overnight. ZIF-8 and ZIF-8^{Pt} were added to the cells and incubated in the dark for 2 h. Following incubation, the medium was removed and cells were washed with PBS three times. Following washing, the ROS probe, DCFH-DA, was added to the cells and incubated for a further 30 min, and then the fluorescence of cells generated by DCFH-DA was detected by fluorescence microscopy.

Cellular Uptake Analysis

HCC cells were seeded in confocal chambers and cultured overnight. The AP loaded Rhodamine B to track nanoparticles, were added to the chambers and incubated for 2 h. Next, the cells were washed with PBS three times and stained with phalloidin-FITC (30 min) and DAPI (10 min). Finally, the cells were observed by laser confocal microscopy (phalloidin-FITC, excitation: 490 nm; DAPI, excitation:400 nm and Rhodamine B, excitation:555 nm). We further investigated the cellular uptake of nanoparticles by FE-TEM. Briefly, HCC cells were cultured in a cell slide overnight. After incubating with $AP@ZIF-8^{Pt}$ for 2 h, the cell slides were washed with PBS. The cells were fixed with 3% glutaraldehyde overnight and then dehydrated in an ethanol series. Using a mini-gold sputter, cell samples were coated with gold-palladium for visual inspection by TEM.

Flow Cytometry Analysis

HCC cells at a concentration of 1×10^5 cells were seeded in six-well plates and incubated overnight. Cells were treated with AP@ZIF-8^{Pt}, which were loaded with rose red B. After incubation for 3 h and 6 h, the cells were washed three times with cold PBS and the treated cells were harvested for flow cytometry analysis. The cells were seeded in 6 well plates at a density of 2×10^5 cells and cultured overnight. Apoptosis kit (BD Biosciences, CA) was used for apoptosis analysis and operated according to the instruction.

Immunofluorescence

HCC cells were fixed with 4% PFA at 37°C for 20 min, treated with 0.02% Triton X-100 for 10 min and then blocked at room temperature for 1 h. Cells were incubated with primary antibody overnight at 4°C followed by incubation with fluorescent-conjugated secondary antibody for 2 h at room temperature. The nuclei were stained with DAPI for 15 min. The images of the cells were obtained by laser confocal microscopy (γ -H2AX-FITC, excitation: 490 nm, HIF-1 α -CY5, excitation: 650 nm and DAPI, excitation: 400 nm).

In vivo Therapeutic Effect

When the tumor reached approximately 100 mm³, tumor-bearing mice were randomly divided into 4 groups: (I) Control; (II) AP; (III) ZIF-8^{Pt}, and (IV) AP@ZIF-8^{Pt}. After the nanoparticles (50 mg·kg⁻¹) were intravenously injected into the tail veins every 2 d. For treatment, the solid tumor was dissected from one mouse in each group for H&E and TUNEL staining. The solid tumor was also used for HIF-1 α and γ -H2AX IHC staining. The body weights of mice and the tumor volumes were recorded every other day.

Transcriptomics Analysis

When the tumor reached about 100 mm³, tumor-bearing mice were randomly divided into three groups: (I) Control; (II) ZIF-8^{Pt}; (III) AP@ZIF-8^{Pt}. Nanoparticles (50 mg/kg) were intravenously injected via tail vein every 2d. At the end of the experiment, tumors were dissected for transcriptomics analysis using an LC-ESI-MS/MS system.

Statistical Analyses

One-way and two-way analyses of variance (ANOVA) and *t*-tests were used to determine statistical significance (*p<0.05, **p<0.01, ***p<0.001), p-value less than 0.05 was considered statistically significant.

Results and Discussion

AP@ZIF-8^{Pt} Synthesis and Characterization

First, zinc nitrate and 2-methylimidazole were used to synthesize zeolitic imidazolate framework 8 (ZIF-8).²⁷ Second, polyvinylpyrrolidone (PVP)-modified Pt nanoparticles was added in to synthesize ZIF-8^{Pt}.^{25,28,29} Third, loaded AP to form AP@ZIF-8^{Pt}. The form of ZIF-8 was first scanned by scanning electron microscope (SEM) (Figure 1A), made sure the framework was correct. The particle size and morphology of AP@ZIF-8^{Pt} was scanned by SEM (Figure 1B) and transmission electron microscopy (TEM) (Figure 1C), verified nanoparticles successful formation. The hydrated particle size of AP@ZIF-8^{Pt} was measured by dynamic light scattering (DLS), which was about 150 nm (Figure 1D). Additionally, the X-ray diffraction (XRD) was performed to analyze the structure of AP@ZIF-8^{Pt}. The result showed that AP@ZIF-8^{Pt} XRD pattern was highly consistent with AP@ZIF-8 alone (Figure 1E), and the peaks (111), (200), (220), and (310) at high angles of 30°–90° corresponding to that of Pt (Figure 1F), suggesting that Pt were coated on ZIF-8 successfully. Further pH-responsive decomposition behavior of AP@ZIF-8^{Pt} was investigated, as shown in Figure 1G, when the pH decreased to 6.0, the accumulative release of AP was significantly increased, suggesting that AP@ZIF-8^{Pt} was easier to degrade in acidic tumor microenvironment. Furthermore, the surface potential of AP@ZIF-8^{Pt} was +26.2 mV measured by Zeta potential (Figure 1H). Finally, size and polydispersity index (PDI) of AP@ZIF-8^{Pt} proved the outstanding stability of AP@ZIF-8^{Pt}, which contained stability in phosphate-buffered saline (PBS) with 10% fetal bovine serum (FBS) for a week (Figure 1I), suggesting that AP@ZIF-8^{Pt} could be used in in vivo potentially.

Cellular Uptake and ROS Generation of AP@ZIF-8^{Pt}

To verify the cellular uptake ability of AP@ZIF-8^{Pt} in HCC cells, AP@ZIF-8^{Pt} was added to Hep3B cells for 6 h. Confocal imaging showed that after 3 h, part of AP@ZIF-8^{Pt} entered the cells and almost all AP@ZIF-8^{Pt} entered the cells after 6 h (Figure 2A), consistently with the flow cytometry analysis for quantitative detection result (Figure 2B). Further field emission transmission electron microscopy (FE-TEM) was applied to verify the real-time absorption of AP@ZIF-8^{Pt} to Huh7 cells. The versatility of FE-TEM allows detailed characterization of the endocytosis of nanoparticles. FE-TEM images showed much AP@ZIF-8^{Pt} entered in HCT-116 cells (marked by blue circle) (Figure 2C). The biocompatibility of ZIF-8^{Pt} was evaluated by Cell Counting Kit-8 (CCK-8) assay, when the concentration of ZIF-8^{Pt} was up to 120 µg/mL, the survival rates of both normal liver cells (LO2) and HCC cells (Hep3B and Hepg2) were more than 95%, suggesting that the superior security of ZIF-8^{Pt} (Figure 2D). The reactive oxygen species (ROS) generation ability of AP@ZIF-8^{Pt} was evaluated by DCFH-DA, an ROS probe to detect H₂O₂ content and oxidative stress.³⁰ As shown in Figure 2E, the green dots had no significant difference in the control and ZIF-8 treatment group, because the endogenous ROS producing in live cells. When treated with ZIF-8^{Pt}, the green dots significantly decreased, as proved in our previous study, Pt nanoparticles could catalyze H₂O₂ to O₂, elevating the source of ROS generation.²⁵ The results showed that ZIF-8^{Pt} presented significant concentration-dependent catalase-like activity.^{31,32}



Figure I Characterization of biomimetic nanoplatform AP@ZIF-8^{Pt}. (A) Scanning electron microscope (SEM) image of ZIF-8. (B) SEM image of AP@ZIF-8^{Pt}. (C) Transmission electron microscopy (TEM) image of AP@ZIF-8^{Pt}. (D) Hydrated particle size of AP@ZIF-8^{Pt}. (E) XRD patterns of AP@ZIF-8 and AP@ZIF-8^{Pt}. (F) XRD patterns in Pt region. (G) The cumulative release profile of AP (n=5). (H) Zeta potential of AP@ZIF-8^{Pt}. (I) Changes of diameter and PDI of AP@ZIF-8^{Pt} over one week.

Therapeutic Efficacy and Mechanism of AP@ZIF-8^{Pt}

Both the human HCC cell lines (Hepg2, Hep3B and Huh7) and the human liver cell line (LO2) were obtained from The American Type Culture Collection (Manassas, VA, USA). CCK-8 assay was first used to verify therapeutic efficacy of AP@ZIF-8^{Pt} against three HCC cell lines (Hepg2, Hep3B and Huh7). The results showed that treatment with AP@ZIF-8^{Pt} significantly inhibits the viability of HCC cells compared with single AP group (Figure 3A). Flow cytometry was then used to verify therapeutic efficacy of AP@ZIF-8^{Pt}. The results showed that the apoptotic rate of AP@ZIF-8^{Pt} treatment group was up to 33.42% for 6 h higher than 3 h (15.48%) (Figure 3B and C). Further therapeutic efficacy of AP@ZIF-8^{Pt} was determined in vivo. Hepg2 tumor-bearing mice were randomly divided into control group, ZIF-8^{Pt} group, AP group and AP@ZIF-8^{Pt} (Figure 3D). The body weights, tumor volumes and tumor weights of the mice were recorded every 2 days in order to assess the effects of different treatments. The body weights of mice increased steadily, demonstrating the biosafety of AP@ZIF-8^{Pt} (Figure 3D). The tumor weight and tumor volumes of mice treated with AP were smaller than those of the control group and ZIF-8^{Pt} group, and the smallest tumors were detected in AP@ZIF-8^{Pt} mice (Figure 3E and F).



Figure 2 Cellular uptake and ROS generation of AP@ZIF-8^{Pt}. (A) AP@ZIF-8^{Pt} incubated with HCC cells for 3 h and 6 h. (B) Flow cytometry analysis for quantitative detection of intracellular AP@ZIF-8^{Pt} fluorescence. (C) Detailed characterization of endocytosis by FE-TEM observation. (D) Cell viability of LO2, Hep3B and Hepg2 cells with AP@ZIF-8^{Pt} treatment were evaluated by CCK-8 assay (n=5). (E) ROS production detected by fluorescence of DCFH-DA in HCC cells.



Figure 3 Therapeutic efficacy and mechanism of AP@ZIF.8^{Pt}. (A) Cell Counting Kit-8 (CCK8) assay results. (B) Apoptosis analysis results. (C) Quantification results of B, n=3. (D) Body weights profiles, (n=5). (E) Tumor weights in each group were measured at the end of the experiment (n=5). (F) Tumor growth profiles (n=5). (G) Histologic analysis and TUNEL staining of tumor sections from different groups of mice (n=5). (H) Immunofluorescence analysis of γ -H2AX and HIF-1 α in tumor tissues collected from various groups of mice (n=5). (*p<0.05, **p<0.01, *** p<0.01).

Moreover, H&E and TUNEL staining analyses were performed to confirm the therapeutic efficacy of AP@ZIF-8^{Pt}. The necrosis and green dots detected in the AP are more than those observed in the control group and ZIF-8^{Pt} group, and the maximum necrosis and green dots were detected in AP@ZIF-8^{Pt} group, indicates the remarkable therapeutic efficacy of AP@ZIF-8^{Pt} (Figure 3G). The mechanism underlying the efficacy of AP@ZIF-8^{Pt} was determined by immunofluorescence staining. The results showed that AP@ZIF-8^{Pt} treatment up-regulated the expression of γ -H2AX, the DNA DSB marker³³ (Figure 3H), indicated that AP@ZIF-8^{Pt} promoted DNA damage by increasing the number of DSBs. In contrast, AP@ZIF-8^{Pt} treatment down-regulated the expression of HIF-1 α , a regulator of primary adaptive responses to hypoxia^{34,35} (Figure 3H), indicated that AP@ZIF-8^{Pt} could alleviate hypoxia in tumor tissue to improve therapy sensitivity.

Transcriptomics Study

Transcriptomics is an extremely important tool for studying all RNA molecules in an organism.³⁶ Cancer-associated protein expression can change levels of some RNA to promote cancer initiation and progression.³⁷ After confirming the in vivo anticancer efficacy of AP@ZIF-8^{Pt}, we further investigated the transcriptomics influence of it in Hepg2-tumorbearing mice. The serum samples from mice treated with PBS, ZIF-8^{Pt} and AP@ZIF-8^{Pt} were used for transcriptomics analysis. The sample density result was shown in Figure 4A. All of the samples in each group were closely clustered, which demonstrated a good quality of samples with different groups. To focus on the differential genes influenced by AP@ZIF-8^{Pt}, we performed analysis of differential genes between two of each of the groups (control and ZIF-8^{Pt}, control and AP@ZIF-8^{Pt} groups and between the control and AP@ZIF-8^{Pt} groups was carried out to eliminate the batch effect. A total of 173 genes were identified between the control and ZIF-8^{Pt} groups, 1223 genes were identified between the ZIF-8^{Pt} and AP@ZIF-8^{Pt} groups. Finally, 1188 genes were identified as significant genes influenced by AP@ZIF-8^{Pt} (Figure 4C).

The enrichment analysis was then performed for these significant genes. As shown in Figure 4D, the top 5 important Kyoto Encyclopedia of Genes and Genomes (KEGG) signaling pathways were thermogenesis, signaling pathways regulating pluripotency of stem cells, ribosome, rion disease and PI3K-Akt signaling pathway. As shown in Figure 4E, the top 3 important biological processes were purine nucleoside diphosphate metabolic process, purine ribonucleoside diphosphate metabolic process and ribonucleoside diphosphate metabolic process. The top 3 important cellular components were respiratory chain, cytosolic large ribosomal subunit and respiratory chain complex. And the 2 important molecular functions were extracellular mata structin consultent and Oxygen binding. Peixoto et al proved that HCC induced by hepatocyte Pten deletion reduced thermogenic capacity.³⁸ As is known to all, the dysregulation of the PI3K-AKT-mTOR signaling pathway is common in HCC.³⁹ Xue et al proved METTL16 promoted liver cancer stem cell self-renewal via controlling ribosome biogenesis and mRNA translation.⁴⁰ Read et al demonstrated that Soluble enzyme hydrolyzes purine nucleoside diphosphate resulted in HCC.⁴¹

Biocompatibility of AP@ZIF-8^{Pt}

It is essential to evaluate the biosafety to determine whether AP@ZIF-8^{Pt} can be applied in clinical treatment.⁴² Considering that changes in the behavior and body weights of mice are considered signs of acute toxicity and that changes in gross pathology reflect long-term toxicity,⁴³ these parameters were assessed throughout the treatment period. Blood samples were collected from mice administered with i.p. injections of PBS, ZIF-8^{Pt} and AP@ZIF-8^{Pt}, and the behavior and body weights of these mice were monitored. The results showed that treatment with ZIF-8^{Pt} and AP@ZIF-8^{Pt} did not induce significant variations in mouse behavior and body weight compared to the control group (Figure 3D), which proved that the acute toxicity of ZIF-8^{Pt} and AP@ZIF-8^{Pt} was negligible. Similarly, routine blood tests proved that the ZIF-8^{Pt} and AP@ZIF-8^{Pt} and AP@



Figure 4 Transcriptomics analysis. (A) Sample density distribution. (B) Volcano map of differential genes in each treatment group. (C) Venn diagrams of significant differential genes influenced by each treatment group. (D) Pathway enrichment analysis of significant genes influenced by AP@ZIF-8^{Pt}. (E) Gene ontology (GO) enrichment analysis of significant genes influenced by AP@ZIF-8^{Pt}.



Figure 5 Biocompatibility of AP@ZIF-8Pt in vivo. (A) Changes of routine blood test contained hemoglobin (HGB), platelet (PLT), red blood cell (RBC) and white blood cell (WBC). (B) Changes of hepatorenal function contained alanine aminotransferase (ALT), aspartate aminotransferase (AST), blood urea nitrogen (BUN) and creatinine (CR). (C) HE staining of mice after three injections of PBS, ZIF-8^{Pt} and AP@ZIF-8^{Pt}.

Conclusion

In this study, a pH-responsive and catalase-like Pt nanoparticle-embedded metal-organic framework (ZIF-8^{Pt}) loaded with AP (AP@ZIF-8^{Pt}) was developed to treating HCC. Both in vitro and in vivo experiment results proved that AP@ZIF-8^{Pt} had excellent biocompatibility and therapeutic efficacy. Further experiment verified that AP@ZIF-8^{Pt} can promote DNA damage and relieving hypoxia. Transcriptomics analysis demonstrated that AP@ZIF-8^{Pt} worked through thermogenesis, signaling pathways regulating pluripotency of stem cells, ribosome, rion disease, PI3K-Akt signaling pathway, purine nucleoside diphosphate metabolic process, purine ribonucleoside diphosphate metabolic process and

ribonucleoside diphosphate metabolic process, respiratory chain, cytosolic large ribosomal subunit, respiratory chain complex, extracellular mata structin consultent and oxygen binding. Our study suggested that this nanoplatform $AP@ZIF-8^{Pt}$ offered a promising paradigm for HCC treatment and had clinical application potential.

Ethics and Consent to Participate Declarations

All animal experiments were approved by the Experimental Animal Management and Ethics Committee of Xiamen University. All experiments were performed in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals.

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Disclosure

This paper has been uploaded to ResearchSquare as a preprint: <u>https://www.researchgate.net/publication/387685173_A</u> <u>super-assembled_synergistically_nanoplatform_APZIF-8_Pt_for_hepatocarcinoma_therapy</u>. The authors declare no competing financial interest.

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