



Original Article

# PAK4-relevant proliferation reduced by cell autophagy via p53/mTOR/p-AKT signaling

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**Background:** P21-activated kinase 4 (PAK4) involves in cell proliferation in cancer and mutually regulates with p53, a molecule is demonstrated to control cell autophagy by mammalian target of rapamycin (mTOR)/protein kinase B (AKT) signaling. Since the signaling exhibits an association with PAK family members in cell autophagy, it implies that PAK4-relevant proliferation may be impacted by autophagy via p53 with a lack of evidence in cancer cells.

**Methods:** In this research, transient and stable PAK4-knockdown human hepatocarcinoma cell lines (HepG2) were constructed by transfection of PAK4-RNA interference (RNAi) plasmid and lentivirus containing PAK4-RNAi plasmid, respectively. We investigated cell proliferation using methyl thiazolyl tetrazolium (MTT) and Cell Counting Kit 8 (CCK8) assays, cell cycle by flow cytometry (FCM) and cell autophagy by monodansylcadaverine (MDC) staining and autophagic biomarker's expression, and detected the expressions of p53, mTOR, phosphorylated-AKT (p-AKT) and AKT by immunofluorescence and western blot to explore the mechanism.

**Results:** We successfully constructed transient and stable PAK4-knockdown HepG2 cell lines, and detected dysfunction of the cells' proliferation. An increased expression of p53, as a molecule of cell-cycle-surveillance on G1/S phase, was demonstrated in the cells although the cell cycle blocked at G2/M. And then, we detected increased autophagosome and autophagic biomarker LC3-II, and decreased expressions in p-AKT and mTOR.

**Conclusions:** The proliferation is reduced in PAK4-knockdown HepG2 cells, which is relative to not only cell cycle arrest but also cell autophagy, and p53/mTOR/p-AKT signaling involves in the cell progress. The findings provide a new mechanism on PAK4 block in cancer therapy.

**Keywords:** P21-activated kinase 4 (PAK4); p53; mammalian target of rapamycin (mTOR); autophagy; cell cycle

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## Introduction

The p21-activated kinases (PAKs) are serine/threonine kinases and usually classified into Group I (PAK1–3) and II (PAK4–6). PAK4 involves in a series of malignant phenotypes such as abnormal growth, cancer stem cells, epithelial-mesenchymal

transition, and drug-resistance (1-4). Its inhibitors are shown in recent research to interfere with the progression of carcinoma cell cycle. The inhibitor KY04031 induces cell cycle arrest in A549 at G1/S phase (5). KPT-9274, a dual PAK4/nicotinamide phosphoribosyl transferase (NAMPT)

inhibitor, fosters reduction of G2/M transition and induction of apoptosis in renal cancer cells (6). An overexpressed orphan gene *INKA1*, interfering with nuclear localization of PAK4, increases the proportion of leukemia stem cells in G0 by in acute myeloid leukemia (7). Silencing PAK4 by small interfering RNA (siRNA) induces a G2/M block in cancer cells because of the dysfunction of Ran GTPase and polo-like kinase 1 as its substrates (8,9). These observations imply that PAK4 modulates cell cycle in multiple phases for cancer cell growth.

p53, a molecule upstream to PAK4 by interacting with p21 (10), protects cells via cell-cycle-surveillance on G1/S phase (11). Therefore, p53-mediated G1/S arrest is speculated to impair proliferation in PAK4-blockdown cancer cells. However, like PAK4, p53-p21 pathway controls all checkpoints from DNA synthesis to cytokinesis including G1/S, G2/M, and spindle assembly during the cell cycle (11-13). Further, there is a mutual regulation between p53 and PAK4 because PAK4 is detected to phosphorylate directly p53, which not only attenuates p53 in transcriptional transactivation activity but also inhibits p53-mediated suppression of hepatocellular carcinoma (HCC) cell invasion (14). Therefore, these pieces of evidence make it easy to understand there remains mechanism underlying PAK4-relevant cell proliferation modulated by p53 besides the process of cell cycle.

Indeed, PAK-relative factors induced cancer cell cycle arrest but also the autophagy following being coupled to acetyltransferases (15). PAK inhibitors in meningioma cells induce cell death with increased autophagy-related protein

levels and chromosomes-markers for mitotic catastrophe (16). The inhibition of potential PAK1 interactors revealed the importance of genes related to the autophagy in PAK1-mediated glioma cell proliferation (17). The autophagy is induced in neuroblastoma by targeting circRNA-195 to downregulate PAK2 expression (18). PAK1-knockout fuels autonomous autophagy under nutrient-limiting conditions in prostate cancer cells (19). It implies that a few PAK family members involve in cell autophagy besides of cell cycle although there is no evidence to connect PAK4 with autophagy.

Furthermore, the autophagy is performed by mammalian target of rapamycin (mTOR)/protein kinase B (AKT) signaling in a few studies. The activator of mTOR increases the expression of PAK1, p-PAK1 and decreases the ratio of LC3B2/LC3B1 in prostate cancer cells (20). Ivermectin-induced autophagy is accompanied with decreased PAK1 expression in breast cancer via the ubiquitination-mediated degradation pathway, which decreases the phosphorylation level of AKT, resulting in the blockade of the mTOR/AKT signaling pathway (21). The ubiquitination-mediated degradation of PAK1 blocks mTOR/AKT1 signaling pathway, leading to increased autophagic flux to treat lung cancer (22). In particular, Atg7 (autophagy gene)-deficient hepatocytes fail to repopulate the liver accompanied with downregulated PAK4 protein in a hepatic injury model, which is overcome by pharmacological mTOR inhibition (23).

Also, p53 is suggested to control cell autophagy via mTOR/AKT signaling. The p53-inhibited mTOR activates autophagy in cultured cells under stress (24) similar as rapamycin-suppressed mTOR (25), while the deficiency of p53 prevents fibroblast autophagy and noticeably increases the activity of mTOR in tumors (25,26). The suppressed autophagy with low expression of autophagic markers presents a decrease in mTOR's kinase activity, i.e., insufficient phosphorylation for AKT in fibroblasts (25). Analogously, the impaired mTOR/AKT signaling induces autophagy in HeLa cells (27). The mTOR/AKT signaling similarly performs to autophagic flux in maintenance of stemness and differentiation potential of P19 embryonic carcinoma cells (28).

According to these observations, a simultaneous modulation is inferred for PAK4-relevant proliferation by autophagy and cell cycle, and mechanistically, p53, mTOR and AKT may involve in the modulation. We present the following article in accordance with the MDAR reporting checklist (available at <https://tcr.amegroups.com/article/view/10.21037/tcr-22-2272/rc>).

### Highlight box

#### Key findings

- PAK4 downregulation is association with autophagy in HepG2 cells.

#### What is known and what is new?

- The proliferation is reduced by cell cycle arrest in PAK4-knockdown HepG2 cells.
- The proliferation is reduced by cell autophagy in PAK4-knockdown HepG2 cells.

#### What is the implication, and what should change now?

- PAK4 involves in cell autophagy besides of cell cycle. Cell autophagy and relative signaling molecules could be considered as biomarkers for PAK4 block in cancer therapy.

## Methods

### *Cell culture and reagents*

HepG2 (human hepatocarcinoma cell line) cells were obtained from American Type Culture Collection (ATCC). The cells were maintained in Dulbecco's Modified Eagle's Medium (DMEM) medium, which is supplemented with 10% fetal calf serum (FCS; Invitrogen, CA, USA), penicillin G (100 U/mL), and streptomycin (100 mg/mL). The cells of HepG2 were cultured in a 37 °C humidified incubator with 5% CO<sub>2</sub> and periodically cultured with 0.25% weight/volume (w/v) trypsin solution.

### *Transient and stable PAK4-knockdown HepG2 cell lines*

PAK4-RNAi were cloned into hU6-MCS-CMV-GFP-SV40-Neomycin vector and sequenced by Genechem Co.,Ltd (Shanghai, China). The shRNA-PAK4 sequence was 5'-GGATGAACGAGGAGCAGAT-3' and 5'-CTTCATCAAGATTGGCGAG-3', and shRNA-Control sequence was 5'-TTCTCCGAACGTGTCACGT-3'. The lentivirus containing PAK4-RNAi plasmid was acquired from Genechem Co.,Ltd. The sequences were same as above.

We transfected commercial PAK4-RNAi plasmid and lentivirus into HepG2 cells in a 24-well plate respectively, and selected steadily infected monoclonal cells with puromycin (2 µg/mL). Lentivirus infection ratio was continuously monitored through immunofluorescence (lentivirus labelled with green fluorescent protein; GFP) and confirmed by western blot.

### *Cell Counting Kit 8 (CCK-8) assay*

As cell proliferation assay,  $2 \times 10^4$  cells of 100 µL were seeded into each well in 96-well plates to culture for 24 h at 37 °C. And the plates were incubated for 10 min at 37 °C after the 10 µL solution was added to each well. The optical density (OD) of each well was detected at 450 nm (Dojindo, Rockville, MD, USA).

### *Methyl thiazolyl tetrazolium (MTT) assay*

The cell viability of HepG2 was evaluated with MTT assay. Briefly, HepG2 cells were digested and plated onto 96-well plates with 85% confluence. After 8 hours, we added 0.01 mL MTT of 5 mg/mL in PBS (phosphate buffered saline) solution, placed the plates in a brood chamber at 37 °C for 4 h, and replaced medium with 0.1 mL DMSO

(dimethylsulfoxide). After incubation at 37 °C for 15 min, the OD of each group was measured at 570 nm using spectrophotometer (Bio-Rad, Hercules, CA, USA).

### *Cell cycle analysis*

The trypsinized cells in suspension ( $2 \times 10^6$ /mL) was fixed with 70% ice-cold ethanol for 12 h at 4 °C. We incubated HepG2 cells in cold PBS containing RNase A (Takara, Japan) for 1 h at 37 °C. Cells were treated with propidium iodide (PI) at a final concentration of 5 µg/mL (Beyotime Biotechnology, Shanghai, China), and incubated at room temperature for 25 min. Finally, cells were subjected to flow cytometry (FCM) and analyzed with Mod Fit LT (Verity Software House, ME, USA).

### *Immunofluorescence*

Before cell seeding, a small 8 mm × 8 mm slide was placed in every well in a 24-well plate. When HepG2 cells approach approximately 60% confluence, cell slides were washed in PBS and fixed in 95% ethylalcohol for 30 min at room temperature. The cell slides were then incubated with primary antibody for 12 h at 4 °C, followed by a brief wash thrice with PBS for 5 min and 1 h incubation at 37 °C with secondary fluorescent antibodies, IgG/Cy3 (bs-0295G-Cy3, Bioss, Beijing) and IgM/RBITC (bs-0368R-RBITC, Bioss, Beijing). Slides were rinsed in PBS for 5 min three times, and nucleus was stained with 1 mg/mL DAPI of 200 µL (Beyotime Biotechnology) for 15 min. After drying, the mounted slides were pictured with a fluorescence microscope. The mean immunofluorescent intensity was calculated for target proteins using the software ImageJ.

### *Western blot*

The cell lysate for immunoblotting was extracted using radio-immuno precipitation assay (RIPA) lysis buffer (Cwbio, Beijing, China) with protease inhibitors (Roche, Germany) added. BCA Assay Kit (Cwbio) was used to quantify protein concentration. Protein sample mixed with a loading buffer (40 µg) was separated on sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) for electrophoresis and transferred in a Tris-Glycine buffer to polyvinylidene fluoride (PVDF) membranes (Millipore, Danvers, USA). The PVDF membranes were blocked by 5% of defatted milk. The membranes were blocked and then incubated with specific primary antibodies for 12 h at 4 °C. After washed thrice with Tris-buffered saline with Tween-20

(TBST), membranes were incubated with secondary antibody for 1 h at 37 °C. Finally, immunoreaction signal was detected with enhanced chemiluminescence (ECL) solution (WBLUC0100, Millipore). Signal intensity of membranes was quantified using ImageJ software (Bio-Rad).

The antibody of anti-human PAK4 was purchased from Santa Cruz, California, USA, anti-Akt, anti-p-Akt, anti-mTOR, anti-LC3, anti-Tubulin, anti-glyceraldehyde-3-phosphate dehydrogenase (GAPDH), anti-p53 from Immuno Way, Texas, USA, and horseradish peroxidase-conjugated secondary antibodies from Gene Tex, Texas, USA.

### *Monodansylcadaverine (MDC) staining*

HepG2 cells were cultured on coverslips in 24-well plates for 48 h as shControl, shPAK4, and positive control were administered by rapamycin groups. After the medium was discarded and cells rinsed with 1× wash buffer, 100 µL MDC (50 µmol/L, Sigma-Aldrich) was added into wells and protected from light for 45 mins. Cells were fixed with 4% paraformaldehyde for 15 min after being washed twice. The images were obtained by fluorescence photometry. Based on the mean density of fluorescence, the statistical analysis was performed among groups.

### *Statistical analysis*

The data were normatively expressed as “mean ± SD” from 3 separate experiments at least. One-way analysis of variance (ANOVA) was used to compare differences among multiple experiment groups, and *t*-test for pairs. Calculations were performed using SPSS 22.0 statistical software. Statistical significance was determined by *P* value less than 0.05.

### *Ethical statement*

The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). Institutional ethical approval and informed consent were waived.

## **Results**

### *Transient and stable knockdown of PAK4 inhibit proliferation in HepG2 cells*

Successful construction of transient PAK4-knockdown HepG2 was verified by western blot ( $P < 0.05$ ; *Figure 1A*). To investigate whether PAK4 was associated with cell proliferation in HepG2 cells, we transfected cells with

PAK4-RNAi plasmid and performed MTT assay by OD value detection on day 1, 2, 3, 4 and 5. We found that there were no statistical differences among group shPAK4, shControl, and Control on day 1 and 2. Meanwhile, the proliferation of cells in group shPAK4 exhibited an obvious reduction compared with group shControl and Control on day 3, 4 and 5 ( $P < 0.05$ ; *Figure 1A*).

Further, we established stable shPAK4 and shControl cell lines using Lentiviral Expression System. HepG2 cells were briefly infected with lentiviral particles containing shPAK4 or scramble sequences. Lentivirus infection ratio was continuously monitored by immunofluorescence (lentivirus labelled with GFP) at hour 48, 72, 96 and 120. Only at hour 96 and 120, the cells showed maximum GFP fluorescence intensity, demonstrating that lentiviral particles were fully integrated into chromosomes of HepG2 (*Figure 1B*).

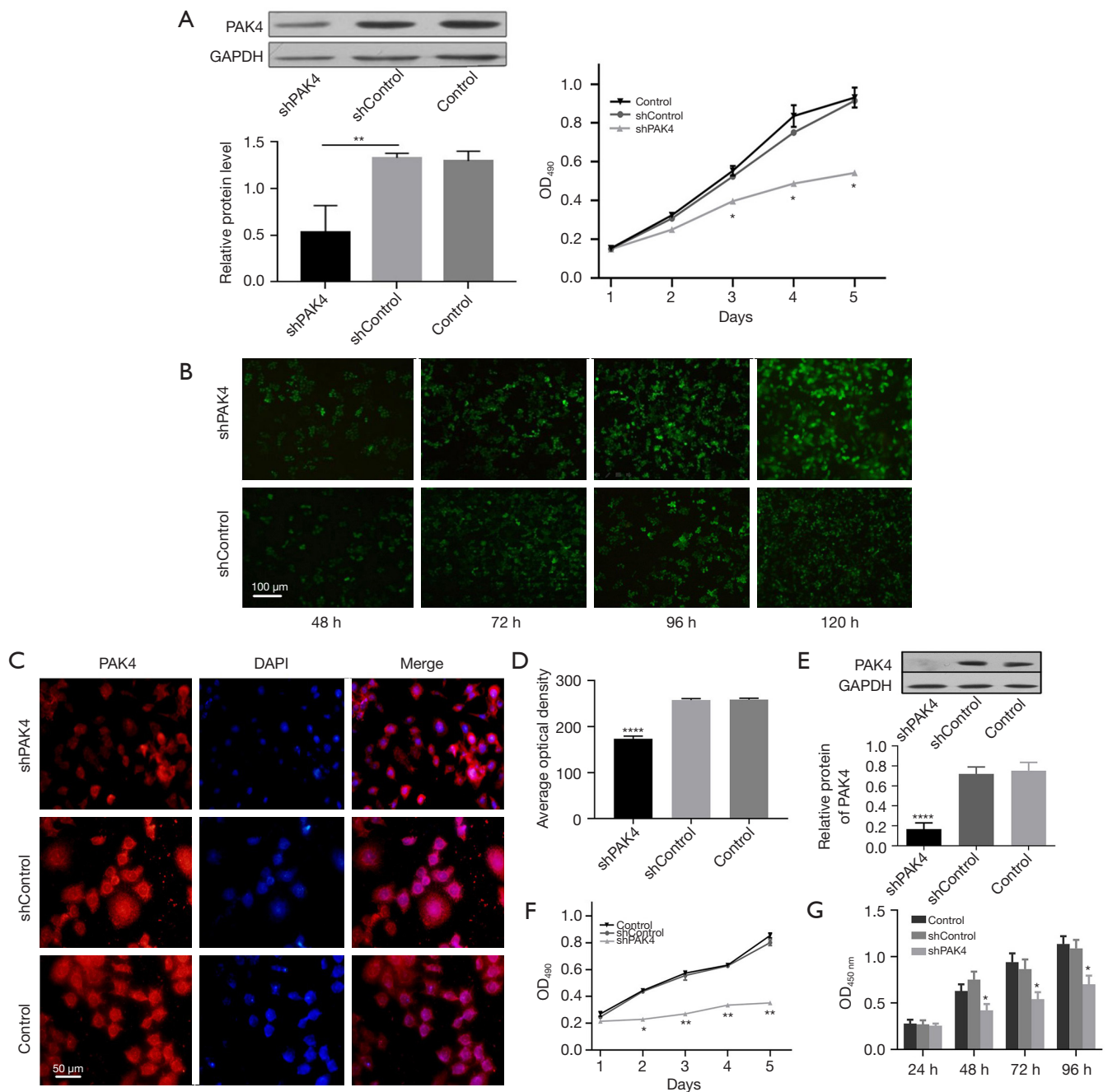
PAK4 was identified for its expression and subcellular localization by immunofluorescence with anti-PAK4 primary antibody and fluorescent-dye conjugated secondary antibody (red). Findings by immunofluorescence showed that PAK4 was expressed in both nucleus and cytoplasm, and there was a reduction in group shPAK4 in comparison to group shControl and Control, which also verified a successful construction of stable shPAK4 cells (*Figure 1C,1D*) as well as western blot (*Figure 1E*). Both results indicated PAK4 was down-regulated in group shPAK4 in comparison to group shControl and Control. Consistent with preceding outcomes, the stable shPAK4 cells showed an attenuated proliferation by MTT or CCK8 assays in comparison to group shControl and Control ( $P < 0.05$  or  $0.01$ ; *Figure 1F,1G*).

### *HepG2 cells are arrested at G2/M phase following knockdown of PAK4*

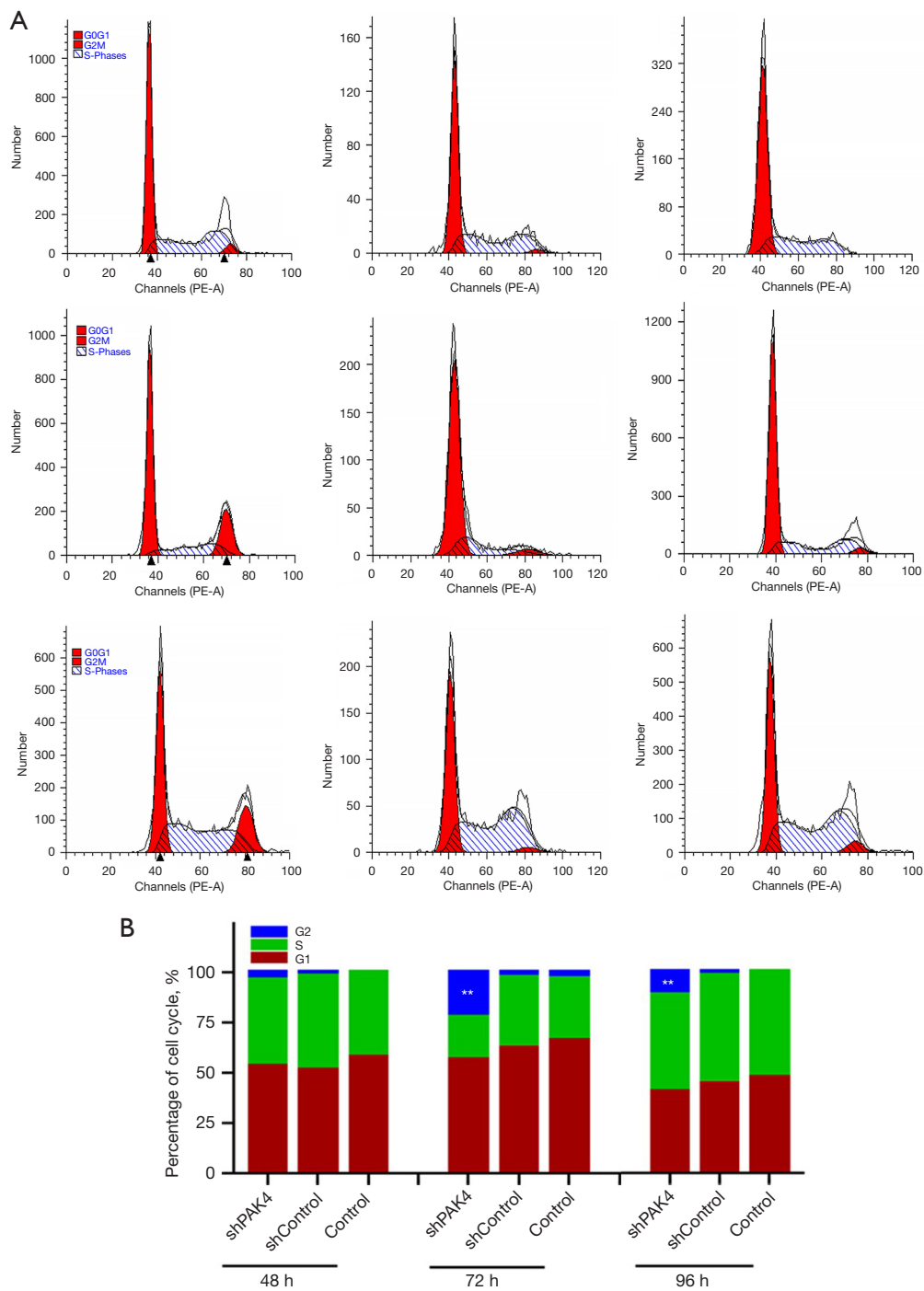
Since the inhibition in proliferation is usually associated with cell cycle arrest, we wonder whether the knockdown of PAK4 results in cell cycle arrest in HepG2 cells. The data by FCM showed that cells exhibited an alteration in cell cycle distribution in group shPAK4, and the proportion of cells was up-regulated at hour 72 and 96 during G2/M phase (*Figure 2A,2B*), indicating knockdown PAK4 induced arrest at G2/M phase.

### *Knockdown PAK4 triggers autophagy phenotype in HepG2 cells*

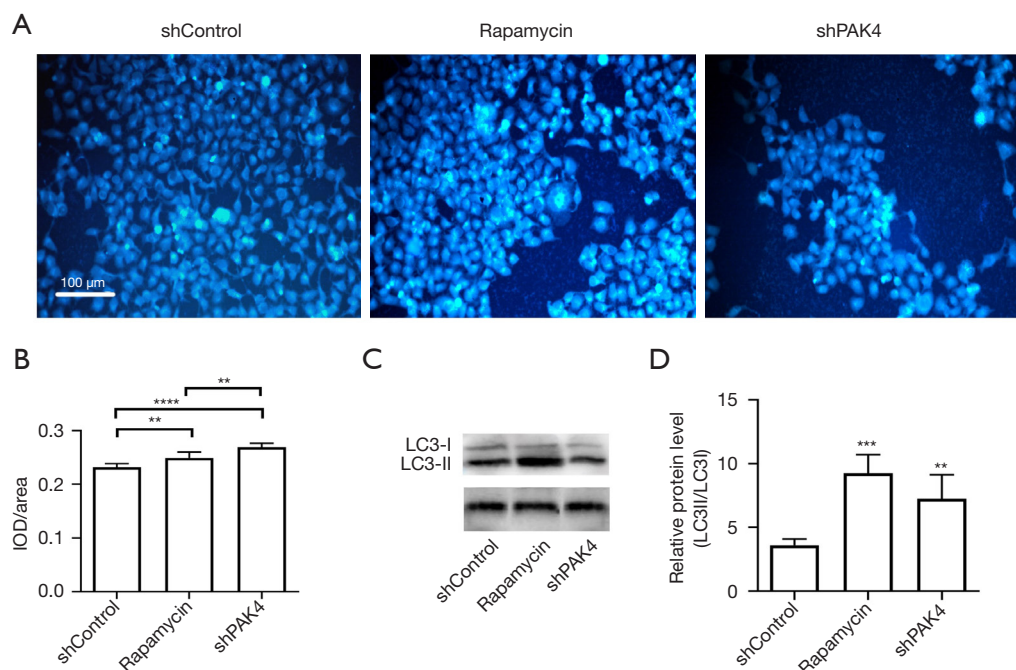
To investigate whether knockdown of PAK4 induces autophagy, autophagosome was checked by MDC staining



**Figure 1** Blockade of PAK4 inhibits proliferation of HepG2 cells. (A) Following transfection with PAK4-RNAi plasmid (shPAK4) or scramble-RNAi plasmid (shControl), PAK4 expression and relative cell viabilities were assessed by western blot and MTT assay. (B) The ratio of HepG2 cells infected with lentivirus was continuously monitored by immunofluorescence (lentivirus labeled with GFP) at hour 48, 72, 96 and 120. (C) The expression of PAK4 were identified by immunofluorescence with anti-PAK4 primary antibody and fluorescent-dye conjugated secondary antibody (red) among groups shPAK4, shControl and Control. DAPI was used to stain nuclei. (D) The fluorescence intensity was statistically analyzed for PAK4. (E) The PAK4 expression levels were verified by western blot. GAPDH was used as an internal control. (F) Following stable knockdown of PAK4 by lentivirus infection, relative cell viabilities were assessed by MTT assay. (G) CCK-8 assay was performed to confirm the effect of PAK4 knockdown on the proliferation in HepG2 cells. \*,  $P < 0.05$ , \*\*,  $P < 0.01$ , \*\*\*\*,  $P < 0.0001$  vs. shControl or Control. PAK4, p21-activated kinase 4; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; OD, optical density; DAPI, 4',6-diamidino-2-phenylindole; HepG2, human hepatocarcinoma cell line; MTT, methyl thiazolyl tetrazolium; GFP, green fluorescent protein; CCK-8, Cell Counting Kit 8.



**Figure 2** Blockade of PAK4 induces HepG2 cells to be arrested at G2/M phase. (A) Cell cycle distribution was detected by flow cytometry among groups shPAK4, shControl, and Control. (B) The proportion of cells at G1, S and G2 phases were analyzed at the 72nd and 96th hour after the knockdown of PAK4. \*\*,  $P < 0.01$  vs. shControl or Control. PAK4, p21-activated kinase 4; HepG2, human hepatocarcinoma cell line.



**Figure 3** Blockade of PAK4 induces autophagy in HepG2 cells. (A) MDC staining was applied to detect autophagosomes in HepG2 cells of each group. (B) The fluorescence intensity of MDC was measured and statistically analyzed among shControl group, Rapamycin group, and shPAK4 group. (C) Western blot analysis was used to detect the conversion of LC3-I to LC3-II. (D) The bar graph displays the gray value ratio of LC3-II to LC3-I. \*\*,  $P < 0.01$ , \*\*\*,  $P < 0.001$ , \*\*\*\*,  $P < 0.0001$  vs. shControl. PAK4, p21-activated kinase 4; IOD, integrated optical density; HepG2, human hepatocarcinoma cell line; MDC, monodansylcadaverine.

in HepG2 cells. As results show, compared with shControl group, there was a larger proportion of fluorescence-positive autophagic vacuoles in group shPAK4 ( $P < 0.01$ ; *Figure 3A,3B*). For further confirmation, western blot was conducted to detect the conversion of LC3-I to LC3-II. Consistent with MDC assay, the expression ratio of LC3-II/LC3-I increased for group shPAK4 ( $P < 0.01$ ; *Figure 3C,3D*). To sum up, there was a negative correlation between PAK4's expression and cell autophagy in HepG2.

#### ***Knockdown PAK4 up-regulates expression in p53 and blockades mTOR/p-AKT signaling***

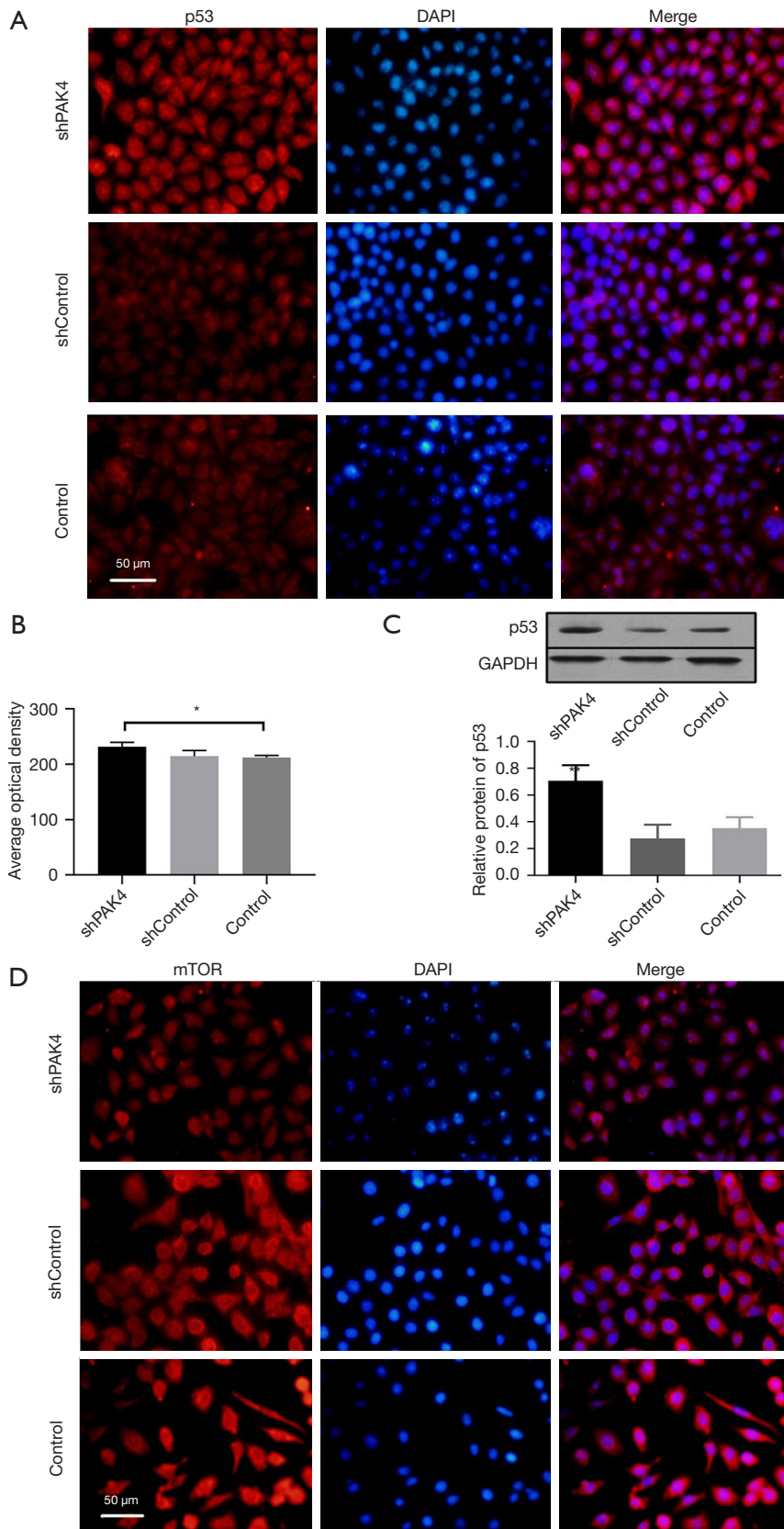
p53 is a pivotal regulator in cell cycle. We detected its expression using immunofluorescence and western blot in PAK4-knockdown HepG2 cells. The findings showed there was a statistically significant up-regulation of p53 in group shPAK4 compared with group shControl and Control (*Figure 4A-4C*), implying that PAK4 negatively regulated p53 in protein. In addition, both mTOR and p-AKT/AKT manifested decreased expressions (*Figure 4D-4H*). It

suggested that high expression of p53 with blocked mTOR/p-AKT signaling in PAK4-knockdown cells.

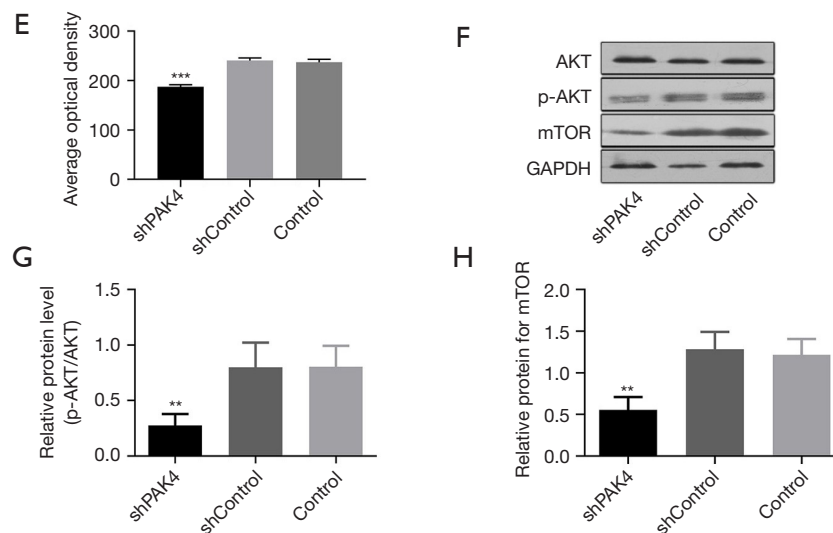
#### **Discussion**

Although p53 is traditionally considered a checkpoint molecule to G1 phase, researches have increasingly revealed that activation of p53 leads to cell cycle arrest at G1/S and G2/M phases (11-13,29). p53-p21 pathway represses cell cycle genes, most of which are essential regulators of G2 phase and mitosis (13). And then, we detected decreased proliferation and cell cycle arrest at G2/M with high expression in p53 in the PAK4-knockdown cells, a phenomenon indicating that the knockdown-induced arrest might be mediated by p53 during G2/M phase in the present study.

Research is rather limited when it comes to the relationship between PAK4 and p53. Although there are reports indicating that PAK4 is upstream to p53, the hierarchy of molecular events is still nubilous during the signal processes connecting PAK4 and p53 (30,31).







**Figure 4** Silenced PAK4 induces upregulation of p53 and blockade of AKT/mTOR pathway. (A) The expression of p53 was measured by immunofluorescence. (B) The fluorescence intensity was statistically analyzed for p53. (C) The expression of p53 was measured by western blot. (D) The expression of mTOR was measured by immunofluorescence. (E) The fluorescence intensity was statistically analyzed for mTOR. (F) The expressions of AKT, p-AKT and mTOR were verified by western blot; GAPDH was used as an internal control mechanism. (G) The gray value ratio of p-AKT to AKT was statistically analyzed. (H) The gray value for mTOR was statistically analyzed. \*,  $P < 0.05$ , \*\*,  $P < 0.01$ , \*\*\*,  $P < 0.001$  vs. shControl or Control. PAK4, p21-activated kinase 4; DAPI, 4',6-diamidino-2-phenylindole; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; mTOR, mammalian target of rapamycin; AKT, protein kinase B; p-AKT, phosphorylated-AKT.

However, we analyzed that PAK4 has a negative regulation to p53 because of their opposite expressions in the study, supported by the inaction of p53 due to its phosphorylation by PAK4 (14). Similarly, a study shows that PAK4 enhances binding between p53 and murine double minute 2, which promotes ubiquitination-mediated p53 degradation and increases activity of glucose-6-phosphate dehydrogenase on pentose phosphate pathway, resulting in an increased multiplication of cells in colon cancer (32).

It is well-known that impaired cell proliferation results from not only cell cycle arrest but enhanced death. Since the death was not from apoptosis in PAK4-knockdown hepatocarcinoma cells in our previous study (33), and p53 (25,26,34) and PAKs (16,19,21) were relevant with autophagy in cancer cells, the present study focused on autophagy. To date, main functional forms in autophagy are proposed to be cytotoxic, cytostatic, and cytoprotective in the context of antitumor therapy: cytotoxic autophagy leads to cancer cell death, cytostatic autophagy inhibits cancer cell propagation, and cytoprotective autophagy enhances cancer cells' drug sensitivity (35). Our findings showed an increased autophagosome by MDC staining and expression

of autophagic biomarker LC3-II/LC3-I, suggesting an increased autophagy inhibiting propagation in PAK4-knockdown HepG2 cells.

Some studies prove that p53-activated cell autophagy, as well as PAKs in tumors (20,21), are regulated via AKT and/or mTOR (25-28,34). Also, mTOR/AKT displays a close relationship with cell autophagy. The inhibition of mTOR/AKT signaling enhances autophagy via facilitating conversion of LC3-I to LC3-II in vascular smooth muscle cells (36). Meanwhile, autophagy is induced along with an inactivation in mTOR/AKT pathway in human breast cancer cells and cervical cancer cells (37,38), while high expression of AKT suppresses autophagy in ovarian cancer (39). In addition, mTOR/AKT is known as a canonical intracellular oncogenic pathway in regulation of tumor cell proliferation (40). Among multiple types of carcinoma, inhibition of mTOR/AKT could enable cell arrests at G2/M phase (41-43). Comparably, our findings exhibited low expressions in mTOR and mTOR's target, p-AKT, suggesting the signaling was associated with the cellular events of cell cycle and autophagy, in PAK4-knockdown HepG2 cells.

## Conclusions

The reduced proliferation in PAK4-knockdown HepG2 cells is caused by cell autophagy, mediated by activated p53 and damp mTOR/p-AKT signaling.

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## Footnote

**Reporting Checklist:** The authors have completed the MDAR reporting checklist. Available at <https://tcr.amegroups.com/article/view/10.21037/tcr-22-2272/rc>

**Data Sharing Statement:** Available at <https://tcr.amegroups.com/article/view/10.21037/tcr-22-2272/dss>

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**Conflicts of Interest:** All authors have completed the ICMJE uniform disclosure form (available at <https://tcr.amegroups.com/article/view/10.21037/tcr-22-2272/coif>). The authors have no conflicts of interest to declare.

**Ethical Statement:** The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). Institutional ethical approval and informed consent were waived.

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