LAB/IN VITRO RESEARCH

e-ISSN 1643-3750 © Med Sci Monit, 2021; 27: e929199 DOI: 10.12659/MSM.929199





MEDICAL

SCIENCE

MONITOR

e929199-1

# Background

Prostate cancer (PCa) has the highest incidence of any male malignant tumor in the United States [1]. Metastasis is among the main features of malignant tumors, leading to a high mortality rate from cancer; therefore, the key factors that initiate this process must be determined.

Speckle-type POZ (pox virus and zinc finger protein) protein (SPOP) is an adaptor protein of E3 ubiquitin ligase that was first identified in the serum of a scleroderma patient by Yasuo Nagai [2]. It is a member of the MATH-BTB protein family; it contains 2 domains: the C-terminal BTB domain and the N-terminal MATH domain. The BTB domain is responsible for binding Cul 3 to form a functional E3 ubiquitin ligase complex, while the function of the MATH domain involves substrate recognition and interaction [3,4]. SPOP substrate research has shown that the BTB protein plays a role in regulating a variety of cellular processes, including hormone-dependent signaling, cell cycle regulation and apoptosis, cell differentiation, and epigenetic control [5-8].

Many studies have shown that SPOP is the most common gene found to be mutated in PCa (10-15%), and that it is concentrated in the MATH-encoding region [9,10]. This suggests that its interaction with substrate proteins may be impaired by mutations. Therefore, some substrates of CRL3<sup>SPOP</sup>, such as the androgen receptor (AR), are the subjects of specific background research in the field of PCa biology [11,12]. Other substrates of CRL3<sup>SPOP</sup> include steroid receptor coactivator 3 (SRC-3) [6], the DEK oncogene [13], and ETS-related gene (ERG) [14-17]. Until now, there has been only a small amount of research on the relationship between metastasis and SPOP. Chen et al reported that SPOP suppressed invasion by osteosarcoma via the PI3K/AKT/NF-kappa B signaling pathway [18]. Xu et al said that the restoration of SPOP could inhibit miRNA-543-induced gastric cancer cell migration and invasion [19]. Duan et al also wrote a preview showing that ERG rearrangements or SPOP mutations lead to elevated ERG levels, thereby promoting cell invasion in PCa [15]. This preview referenced the results of Gan and An, who found that SPOP mutations in PCa inhibit the degradation of ERG, leading to enhanced levels of ERG and the promotion of cell invasion [16,17].

Despite the relevant studies mentioned above, the specific mechanisms through which SPOP mutations promote tumor metastasis are unclear. In recent years, more attention has been paid to the effects of the tumor microenvironment on tumor metastasis [20,21]. In the current study, we have focused on the microenvironment to explain cell metastasis in SPOP-mutant PCa. With a view toward further revealing the molecular mechanism underlying SPOP-mutant PCa, we first obtained SPOP knockout PCa cells and verified the change in their metastasis ability. Furthermore, we found that MCP-1 was highly expressed in SPOP knockout cell-conditioned medium (CM) and that it may be the key molecule that promotes SPOP-mutant PCa metastasis.

# **Material and Methods**

## **Cell Culture**

The SPOP knockout human PCa cells,  $C4-2^{SPOP-/-}$  sgSPOP-1 and sgSPOP-2, and the control cells were gifts of Dr. Wenyi Wei at Harvard Medical School. The cells were cultured in Royal Park Memorial Institute (RPMI 1640) medium (GIBCO, California, USA), containing 10% Fetal Bovine Serum (FBS) and streptomycin, at 37°C in a humid, 5%  $CO_2$  atmosphere. The cells were routinely tested for mycoplasma contamination.

### Preparation of Cell-conditioned Medium

CM from sgSPOP-1, sgSPOP-2, and control cells was harvested for soluble factors, which were measured and subsequently verified by enzyme-linked immunosorbent assay (ELISA) [22].

## Western Blotting

C4-2<sup>SPOP-/-</sup> sgSPOP-1 and sgSPOP-2, as well as control cells, were lysed in a radio immunoprecipitation assay (RIPA) buffer containing a protease inhibitor cocktail. After centrifugation, the supernatant was collected, the concentration was measured, and denaturation was allowed to occur. Then, the whole-cell lysates (50 µg) were resolved through SDS-polyacrylamide gel electrophoresis (SDS-PAGE), and then the proteins were transferred from the gel to a polyvinylidene difluoride (PVDF) membrane and incubated in a blocking buffer that consisted of 5% nonfat milk with tris buffered saline with Tween (TBST) buffer, for 1 h at room temperature. After washing with TBST buffer, the membranes were incubated with a primary antibody to SPOP (Proteintech, Wuhan, China) and tubulin (Cell Signaling Technology, Boston, USA) overnight at 4°C. After another TBST washing, followed by another 1 h of incubation with a secondary HRP antibody, at room temperature, immunoreactive proteins were detected using an enhanced chemiluminescence SuperSignal substrate (Thermo Fisher Scientific, Massachusetts, USA).

### Transwell Invasion Assays and Transwell Migration Assays

Non-FBS medium (500  $\mu$ L) was put into 8.0  $\mu$ m 24-well plate chamber inserts (Corning 3422 for migration and BD Biosciences for invasion, Massachusetts and New Jersey, USA, respective-ly). After removal from the refrigerator, the Matrigel-medium mix was allowed to solidify through incubation at 37°C for 2 h.

Then, the plates were removed from the incubator, and 5×10<sup>4</sup> cells in serum-free medium were seeded in the top chamber. The bottom of each insert was filled with 750 µl of medium supplemented with 10% FBS. For the specific factor blocking assay, RPMI 1640 medium and CM with 2 µg/ml monoclonal mouse anti-human MCP-1 (eBioscience, California, USA) were placed in the lower chambers. For each sample, the procedure was repeated 3 times. After 24 h of cell incubation, the cells were washed with phosphate buffer saline (PBS) and fixed with 4% paraformaldehyde for 18 min. Cotton swabs were used to clear up the cells on the top of the insert, while 0.5% crystal violet blue was used to stain the cells that adhered to the bottom. After 15 min of standing at room temperature while protected from light, they were washed with H<sub>2</sub>O. The images were obtained using a microscope and the cells were counted in 15 different view fields using ImageJ software. An invasive index was used to calculate the ratio of the cells that had invaded through the Matrigel-coated membrane to the cells that migrated through the uncoated membrane [22].

## Wound-healing Assay

C4-2<sup>SPOP-/-</sup> sg-SPOP-1 and sg-SPOP-2 cells and control cells were plated in six-well plates at a high concentration of  $5 \times 10^5$  cells/ well. After 24 h of incubation, the cells were treated with 20 µg/ml mitomycin for 3 h to inhibit proliferation. Scratches were made in the plate, using a 200 µl pipette tip. Images were collected at 0, 10, and 18 h after scratching. The experiment was repeated 3 times. Wound closure areas were quantified using Photoshop and ImageJ software.

# Antibody Array

Human Chemokine Array C1 from RayBiotech (Atlanta, GA, USA) contains 38 repeated antibodies spotted on a single membrane. Experiments were implemented according to the manufacturer's protocol, and the results were analyzed by both Lane 1D software from Sage Creation (Beijing, China) and software from RayBiotech.

### ELISA

To measure the expression/secretion levels of soluble factors, the conditioned media were harvested from C4-2<sup>SPOP-/-</sup> sgSPOP-1 and sgSPOP-2 and control cells. An MCP-1 ELISA kit from NeoBioscience (Shenzhen, China) was used to measure the levels of MCP-1 in the conditioned medium. This was performed according to the manufacturer's protocol.

# Quantitative PCR (qRT-PCR)

TRIzol (Invitrogen, California, USA) was used to extract total RNA from C4-2<sup>SPOP-/-</sup> sgSPOP-1 and sgSPOP-2 cells, and control

cells, according to the manufacturer's instructions. The total RNA concentration was detected by a NanoDrop 2000 spectrophotometer (Thermo Fisher Scientific). A High-Capacity cDNA Reverse Transcription Kit (Invitrogen) was used to convert RNA into cDNA, according to the manufacturer's protocol. The following primers were used:

MCP-1: forward 5'-TAGCAGCCACCTTCATTC-3' and reverse 5'-GGAATCCTGAACCCACTT-3'; human GAPDH: forward 5'-AGCCACATCGCTCAGACA-3' and

reverse 5'-GCCCAATACGACCAAATCCA-3'.

Additionally, qRT-PCR was performed using the ABI 7500 system (Applied Biosystems, California, USA). The experiment was conducted independently in 3 biological replicates, and GAPDH was used as the housekeeping control gene.

## **Statistical Analysis**

StatView software (Abacus Concepts, Berkeley, CA) was used to perform statistical analysis. The significance of the in vitro cell migration, cell invasion, ELISA, and qPCR between the C4-2<sup>SPOP-/-</sup> (sgSPOP-1, sgSPOP-2) and control groups was determined by a *t* test (2-tailed). Data are expressed as mean±standard deviation (SD). All statistical analyses were implemented with SPSS Statistics 16.0. *P*<0.05 was considered statistically significant (\* *P*<0.05, \*\* *P*<0.01).

# Results

### SPOP Knockout Promotes C4-2 Cell Migration And Invasion

Western blotting was used to confirm that sgSPOP-1 and sgSPOP-2 (**Figure 1A**) cells did not express SPOP, but control cells did express SPOP (**Figure 1B**). The results showed that SPOP was knocked out in sgSPOP-1 and sgSPOP-2.

To further investigate the metastasis-related function of SPOP, transwell assays were implemented between sgSPOP-1 and sgSPOP-2 (without SPOP expression) and control cells (with SPOP expression). The results revealed that sgSPOP-1 and sgSPOP-2 cells showed stronger migration and invasion abilities compared with control cells (**Figure 1C**). Subsequently, a scratch wound-healing assay was performed, and it showed that sgSPOP-1 and sgSPOP-2 cells had higher migration abilities than the control cells. This was consistent with the results of the transwell assay (**Figure 1D**).

### C4-2 SPOP<sup>-/-</sup> cells produce excessive amounts of cytokines

C4-2 SPOP<sup>-/-</sup> cells showed stronger migration and invasion abilities than C4-2 control cells. To characterize the changes in soluble factors secreted by C4-2 SPOP<sup>-/-</sup>, CM was collected from sgSPOP-1 and sgSPOP-2 and control cell cultures, and a

Human Chemokine Antibody Array (RayBiotech) was applied to detect the expression of chemokines in these media. After normalization, the results demonstrated that the expression of MCP-1 in sgSPOP-1 and sgSPOP-2 was upregulated compared with that in the control cells (**Figure 2A**).

To verify the antibody array data, ELISA was applied to characterize the secretion levels of MCP-1 in the collected CM. In accordance with the results of the arrays, the expression/secretion levels of MCP-1 in the C4-2 SPOP<sup>-/-</sup> cells were significantly higher than those in the C4-2 control cells (**Figure 2B**). Furthermore, this result was also confirmed by qRT-PCR, in which the expression of MCP-1 was higher in the C4-2 SPOP<sup>-/-</sup> cells (**Figure 2C**). Since MCP-1 in the CM enhanced the migration and invasion of PCa cells, and since MCP-1 was expressed at higher levels in the CM of C4-2 SPOP<sup>-/-</sup> cells, we focused on the association between MCP-1 and C4-2 SPOP<sup>-/-</sup> cell migration and invasion. To test whether MCP-1 in the CM of C4-2 SPOP<sup>-/-</sup> cells affects their migration and invasion, we conducted a transwell assay. Many migrating and invading cells were recognized in the CM-treated group, whereas fewer migrating and invading cells were found in the control medium (**Figure 3**). The CM with the anti-MCP-1 antibody reduced the number of migrating and invading cells (P<0.01 for sgSPOP-1 migration and P<0.05 for sgSPOP-2 migration; P<0.05 for both sgSPOP-1 and sgSPOP-2 invasion).



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Indexed in: [Current Contents/Clinical Medicine] [SCI Expanded] [ISI Alerting System] [ISI Journals Master List] [Index Medicus/MEDLINE] [EMBASE/Excerpta Medica] [Chemical Abstracts/CAS]



Figure 1. SPOP knockout promotes C4-2 cell migration and invasion. (A) Photomicrograph of sgSPOP-1 and sgSPOP-2 compared with parent cell line C4-2 (control). (B) Western blotting with anti-SPOP antibody confirms the presence of SPOP in control C4-2 cells and absence in both sgSPOP-1 and sgSPOP-2 cells. (C) A transwell assay was used to explore cell migration and invasion ability in sgSPOP-1, sgSPOP-2, and control cells. The results are presented as the mean±SD of 3 independent replicates;
\* P<0.05, \*\* P<0.01. (D) The migration ability of sgSPOP-1 cells, sgSPOP-2 cells, and control cells was investigated using a scratch wound-healing assay. The results are shown as the mean±SD of 3 independent replicates; \* P<0.05, \*\* P<0.01.</li>

# Discussion

Among men worldwide, PCa is the type of malignant tumor with the highest incidence. Recurrence and metastasis are the biggest challenges for clinical treatment [23]. In PCa, SPOP is the most common mutated gene, and there are reports showing that SPOP is closely related to tumor cell metastasis [16,17,19,24]. We hypothesized that the loss of SPOP may promote PCa metastasis by changing the tumor microenvironment. Certain protein networks regulate the occurrence of metastasis. Blocking SPOP mutation or the mutation of related key molecules may benefit patients who are at risk of metastasis.

In the current study, we first obtained SPOP knockout PCa cells through a lentiCRISPR/Cas 9 system (C4- $2^{\text{SPOP}-/-}$  and control cells). Then, western blot was used to verify SPOP expression. After determining that the loss of SPOP creates stability, we found that C4- $2^{\text{SPOP}-/-}$  showed stronger migration and invasion abilities compared with control cells, through transwell and wound-healing assays. This result is consistent with the previous results of other scholars [15].

Next, we wanted to know how SPOP suppresses tumor metastasis. Our results suggested that the loss of SPOP changed the microenvironment that promotes tumor cell migration and invasion. By using antibody arrays, we identified that the MCP-1 level dramatically increased in the C4-2<sup>SPOP-/-</sup> cells' CM compared with the control cells' CM. ELISA and qRT-PCR verified this finding. These results show that the loss of SPOP may promote PCa cell migration and invasion through upregulation of MCP-1 in the tumor microenvironment. To confirm this finding, we used an anti-MCP-1 antibody to neutralize MCP-1 in the CM of C4- $2^{\text{SPOP-/-}}$ . A transwell assay was again used to compare the migration and invasion abilities of C4- $2^{\text{SPOP-/-}}$  cells in the following media: RPMI 1640, CM, and CM+anti-MCP-1. C4- $2^{\text{SPOP-/-}}$  cells showed decreased migration and invasion abilities in CM+anti-MCP-1, which suggested that MCP-1 may be a key molecule enabling metastasis in SPOP-mutant PCa.

MCP-1, also known as CCL2, belongs to the C-C chemokine subfamily. It has been shown to be important for recruitment of monocytes, activation of the process of acute inflammation, and development of cancer. Extensive research has indicated that MCP-1 could induce the occurrence of EMT and promote cell proliferation [25-27], angiogenesis [28-31], and metastasis [29,32,33]. Higher levels of MCP-1 after neoadjuvant chemotherapy may lead to recurrence and metastasis [34]. Additionally, in the tumor microenvironment, higher MCP-1 expression may enhance the likelihood of cancer cell invasion by activating both autocrine and paracrine pathways [35]. These findings are consistent with our finding that MCP-1 secreted by SPOP-mutant PCa cells may promote cell migration and invasion.



Figure 2. The difference in soluble factors in the conditioned media of sgSPOP-1 cells, sgSPOP-2 cells, and control cells, as detected by chemokine antibody arrays and verified by ELISA and qRT-PCR. (A) The difference in expressed chemokines in the conditioned media of sgSPOP-1 cells, sgSPOP-2 cells, and control cells, as detected through chemokine antibody array. (B) ELISA was used to verify the MCP-1 expression level in the conditioned medium. The results are presented as the mean±SD of 3 independent experiments; \*\* P<0.01. (C) qRT-PCR measurement of the MCP-1 mRNA expression levels. The results are presented as the mean±SD of 3 independent experiments; \* P<0.05, \*\* P<0.01.</p>

Much research has shown that SPOP is a frequent mutation factor in PCa. Additionally, its structure, cellular function, functional consequences of mutation, and possible mechanisms in PCa have been explored in many reports. An et al and Gan et al reported that cancer-associated SPOP mutations were not sufficient to promote ERG ubiquitylation. Furthermore, they found that ERG stabilization is important for enhancing the migration and invasion activities of SPOP-mutant cells [16,17]. Our research focused on changes in the tumor microenvironment and found that SPOP could regulate the secretion of MCP-1, thereby further regulating cell migration and invasion in PCa. Although SPOP may affect the metastasis of PCa through its substrate, according to the result of the transwell assay with or without anti-MCP-1 antibody, the occurrence of PCa cell migration and invasion is at least partially achieved through the involvement of MCP-1.

Studies have shown that SPOP can mediate the degradation of BRD4 [36,37]. On the one hand, in SPOP-deficient PCa cells, enhanced BRD4 transcription co-activation and increased ERG expression jointly promoted cell migration. On the other hand, BRD4 plays an important role in the production of CCL2. It upregulates the expression of MCP-1 via the activation of the NF- $\kappa$ B signaling pathway. In other words, SPOP regulates MCP-1 in an indirect way. Many phase I and phase II clinical trials have shown the efficacy of the monoclonal antibody carlumab



Figure 3. The migration and invasion of sgSPOP-1 and sgSPOP-2 cells in response to their CM. (A) A total of 5×10<sup>4</sup> sgSPOP-1 cells were seeded to test for migration and invasion ability. Lane 1, RMPI 1640; Lane 2, sgSPOP-1 CM; Lane 3, sgSPOP-1 CM with anti-MCP-1 (2 μg/ml). (B) A total of 5×10<sup>4</sup> sgSPOP-2 cells were seeded to test for migration and invasion ability. Lane 1, RMPI 1640; Lane 2, sgSPOP-2 CM; Lane 3, sgSPOP-2 CM with anti-MCP-1 (2 μg/ml). The results are presented as the mean±SD of 3 independent experiments; \* P<0.05, \*\* P<0.01. CM – conditioned medium.</p>

targeting MCP-1 in solid tumors [36-38]. Prostate cancer patients may also benefit from this approach.

# Conclusions

The loss of SPOP can promote PCa cell migration and invasion. In this study, we explored the possible mechanisms underlying this promotion using antibody arrays and verified array results. However, to better understand the molecular mechanisms behind the association of SPOP and MCP-1 with metastasis, further studies are needed. Additionally, in vivo studies are necessary. MCP-1 and related pathway inhibitors can be used to block the migration and invasion of PCa, thereby preventing tumor recurrence and metastasis.

### Acknowledgements

The authors would like to thank Dr. Xiangpeng Dai for help-ful discussions.

### **Conflict of Interest**

None.

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