

Fascin lysine 471 acetylation cooperates with serine 39 phosphorylation to inhibit actin-bundling activity and tumor metastasis in esophageal squamous cell carcinoma

Dear Editor,

Fascin actin-bundling protein 1 (Fascin, gene name: *FSCNI*) is the primary actin-bundling protein that clusters filaments into stable and bundled filopodia [1]. In previous studies, elevated levels of Fascin promoted filopodia formation and enhanced the invasive ability of tumor cells, suggesting that Fascin is a target for therapeutic intervention [2, 3]. Therefore, it is necessary to elucidate the detailed molecular mechanism by which Fascin bundles F-actin filaments.

Rapid association/dissociation of Fascin from filaments is regulated by its post-translational modifications, such as the widely studied phosphorylation of conserved Ser-39 mediated by protein kinase C (PKC) [4]. The phosphorylation/dephosphorylation cycles of Ser-39 serve as a molecular switch to turn Fascin activity on or off to determine filopodia formation [5]. We previously reported that Fascin was acetylated by P300/CBP-associated factor (PCAF) on Lys-471 [6]. We further elaborated on the precise regulatory mechanism of Fascin acetylation to inhibit its oncogenic function by reducing actin-bundling activity in esophageal squamous cell carcinoma (ESCC) cells, highlighting the

List of abbreviations: A, alanine; AcK471-Fascin, Fascin-K471 acetylation; CK, cytokeratin; D, aspartic acid; DFS, disease-free survival; ESCC, esophageal squamous cell carcinoma; FascinS39A/K471Q, a mimic of phosphorylation defective and hyperacetylated Fascin; FascinS39A/K471R, a mimic of phosphorylation and acetylation defective Fascin; FascinS39D/K471Q, a mimic of hyperphosphorylation and hyperacetylated Fascin; FascinS39D/K471R, a mimic of hyperphosphorylation and acetylation defective Fascin; FRAP, fluorescence recovery after photobleaching; G/F-actin ratio, ratio of G-actin to F-actin; GFP, Green fluorescent protein; H&E, hematoxylin and eosin; IHC, immunohistochemistry; Lys-471, lysine 471; OS, overall survival; P, pellet; PCAF, P300/CBP-associated factor; PhoS39-Fascin, Fascin-S39 phosphorylation; PKC, protein kinase C; PTMs, posttranslational modifications; S, supernatant; Ser-39, Serine 39; SHEEC, Shantou Human Embryonic Esophageal Carcinoma; Thr-403, Threonine 403; TMAs, tissue microarrays; TNM, tumor-node-metastasis; WT, wild-type.

potential therapeutic value of mediating Fascin acetylation in tumor metastasis [6]. The phosphorylation of Ser-39 at the Fascin N-terminus and acetylation of Lys-471 at the Fascin C-terminus were inhibited Fasin F-actin bundling activity. However, whether Lys-471 acetylation cooperates with Ser-39 phosphorylation to regulate Fascin function is still unknown. The purpose of this study was to explore the role of Fascin Ser-39 phosphorylation and Lys-471 acetylation in ESCC cell function and filopodia formation.

In this study, we engineered plasmids encoding double point mutations that mimic all four possible modification states. Fascin^{S39A/K471R} (S39A/K471R, a phosphorylationand acetylation-defective Fascin mutant), Fascin^{S39A/K471Q} (S39A/K471Q, a phosphorylation-defective and constitutively acetylated Fascin mutant), Fascin^{S39D/K471R} (S39D/K471R, a constitutively phosphorylated and acetylation-defective Fascin mutant), and Fascin^{S39D/K471Q} (S39D/K471Q, a constitutively phosphorylated and constitutively acetylated Fascin mutant), and expressed them individually in endogenous FSCN1-silenced cells (Figure 1A). The effect of silencing endogenous Fascin could be rescued by exogenous wild-type (WT) Fascin. Cells expressing Fascin^{S39A/K471R} exhibited stronger oncogenic abilities than those expressing the WT form. In contrast, the expression of $Fascin^{S39A/K471Q}$ or Fascin^{S39D/K471Q} produced the opposite effect, that is, it further suppressed cell migration (Figure 1B) and cell proliferation (Supplementary Figure S1A) in FSCN1silenced cells. These data showed that cell oncogenic abilities could be enhanced or rescued by Fascin^{S39A/K471R} or Fascin^{S39D/K471R}, while could not be rescued by Fascin^{S39A/K471Q} or Fascin^{S39D/K471Q}, suggesting a functional interplay between phosphorylation of Ser-39 and acetylation of Lys-471. The in vivo tumor growth of cells expressing Fascin^{S39A/K471Q} and Fascin^{S39D/K471Q} was significantly inhibited (Supplementary Figure S1B and Figure 1C), whereas the deacetylated mutants (S39A/K471R and S39D/K471R) displayed dramatically increased tumor

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FIGURE 1 Fascin acetylation at Lys-471 and phosphorylation at Ser-39 cooperatively modulate ESCC cell biological behavior. (A) Western blotting after transfection of the GFP vector (Vector), GFP-Fascin (WT), or Fascin double-mutant plasmid in stable *FSCN1*-silenced KYSE150 and SHEEC cells. β -Actin was used as a loading control. (B) Migration of *FSCN1*-silenced KYSE150 (top) and SHEEC (bottom) cells transfected with the GFP vector (Vector), GFP-Fascin (WT), or Fascin double-mutant plasmid was investigated by using an xCELLigence Real-Time Cell Analysis DP instrument. Left panel, real-time detection of impedance response/cell index curves determined over 48 h. Right panel, slope of the cell profiles. Data are presented as the mean \pm SD (n = 3). (C) Representative excised primary tumors (from the footpad to the heel of the foot; upper images) and invaded tumors (from the heel of the foot to the inguinal lymph node; lower images) of mice. Primary tumor weights are compared among the six groups. Data are presented as the mean \pm SD. For Vector, WT, S39D/K471R, S39D/K471Q, and S39A/K471Q mice groups, n = 8; for the S39A/K471R mice group, n = 7. (D) Left panel, representative H&E and IHC staining for GFP to detect the primary tumor (upper images) and representative IHC staining of CK in the inguinal lymph nodes (lower images). Bars, 20 μ m.

growth (Figure 1C). These data are consistent with the incidence of lymph node metastasis in the different groups (Figure 1D), highlighting only dephosphorylated and deacetylated (S39A/K471R) Fascin to be a more prominent player. Any post-translational modification at either site inhibited the ability of Fascin to promote tumor growth.

Next, we investigated the functions of these two post-translational modifications in filopodia formation by performing live imaging of cells co-expressing an F-actin reporter gene (Lifeact-RFP) and GFP-tagged WT or double-mutant Fascin. First, the expression of Fascin^{S39D/K471Q} significantly reduced filopodial density and length in both KYSE150 (Figure 1E and Supplementary Video S1) and SHEEC (Supplementary Figure S2A) cells. In FSCN1-silenced cells, the double-defective mutant (S39A/K471R), but not the other mutants, rescued filopodia formation, with elongated and overabundant filopodia being evident (Figure 1E and Supplementary Video S1). These data suggested that either Lys-471 acetylation or Ser-39 phosphorylation was sufficient for initiating filopodia formation, while maximal formation of filopodia requires the absence of modifications at both Lys-471 and Ser-39 residues. We further analyzed the dynamics of the filopodia extension-retraction cycle in FSCN1-silenced cells following transfection with the double-mutant-expressing plasmid. Time-lapse imaging showed that WT Fascin, but not Fascin^{S39D/K471Q} and Fascin^{S39A/K471Q}, could restore filopodia lifetime (Supplementary Figure S2B). Moreover, expression of the deacetylated and dephosphorylated mutant (S39A/K471R) led to long, persistent filopodia with an increased lifetime (Supplementary Figure S2B). Together, these data showed that filopodia assembled in cells with acetylated Fascin were

significantly shorter, less stable, and collapse rapidly. In contrast, deacetylated Fascin enhanced filopodia stability and length, which were further enhanced by dephosphorylation of Ser-39 and reduced by phosphorylation of Ser-39.

Our previous results showed that Lvs-471 acetvlation increases the mobility of Fascin molecules compared to the WT form [6]. Similarly, rapid mobility was observed in KYSE150 (Figure 1F) and SHEEC (Supplementary Figure S2C) cells expressing Fascin^{S39D/K471Q}. However, for the mutant Fascin forms with at least one deactivating modification (S39A/K471Q, S39D/K471R, and S39A/K471R), Fascin mobility was reduced, particularly for the mutant with both sites deactivated (S39A/K471R), where mobility was completely abolished (Figure 1F and Supplementary Figure S2C). Previous studies have suggested that monomeric proteins move more rapidly than proteins with short transient interactions, whereas proteins with long transient interactions move the slowest [7]. Thus, our data indicated that at the base of the filopodium, Lys-471 acetylation and Ser-39 phosphorylation cause the majority of Fascin molecules to exist as monomers with fast dynamics due to reduced actin binding. We next explored whether the modification of one amino acid affected the actin bundling of the other. Expectedly, the phosphorylation and dephosphorylation of Ser-39 have previously been shown to decrease and enhance, respectively, the actin-bundling properties of Fascin [4]. Consistent with our earlier findings, expression of only WT and Fascin^{S39A/K471R} restored the level of F-actin in KYSE150 and SHEEC cells, as shown by the lower ratio of G/F-actin (Figure 1G and Supplementary Figure S3). In contrast, the other three mutants failed to rescue F-actin levels, but instead increased the G/F-actin ratio compared with that of WT Fascin (Figure 1G and

Right panel, number of mice with and without lymph node metastases 32 days after inoculation. (E) Fascin and F-actin distribution and mean filopodial length and density in KYSE150-shFSCN1 cells co-expressing different types of GFP-Fascin (WT, S39D/K471R, S39D/K471Q, S39A/K471R, or S39A/K471Q) along with Lifeact-RFP (at least 30 cells were measured). (F) FRAP assays were used to analyze the diffusion of different types of GFP-Fascin and Lifeact-RFP at the filopodial base of cells. Whole-cell filopodia images were obtained (left) with time-lapse sequence images of the boxed region in one filopodium, before and after photobleaching, shown to the right. The recovery profile was curve-fit using a double exponential function with recovery constant Kinflow and Koutflow for intra-filopodia exchange and filopodia-cytoplasmic exchange, respectively (low; F; n = 30). (G) F-actin bundling activity of Fascin and its double mutant forms was analyzed by western blotting. The pellet (P) and supernatant (S) represent F-actin and G-actin, respectively. Bar charts show the densitometric G-actin/F-actin ratio. Data were obtained from three independent experiments and are presented as the mean \pm SD. (H) Representative IHC staining images of PhoS39-Fascin and AcK471-Fascin in paired ESCC tissues and adjacent normal tissues (n = 146). For statistical analysis, each protein expression score was divided into 2 subgroups, high-expression and low-expression, on the basis of X-tile software analysis. Low: the expression of PhoS39-Fascin \leq 64 scores and AcK471-Fascin \leq 75 scores; High: the expression of PhoS39-Fascin >64 scores and AcK471-Fascin > 75 scores. Other: the expression of PhoS39-Fascin > 64 scores and AcK471-Fascin \leq 75 scores, the expression of PhoS39-Fascin \leq 64 scores and AcK471-Fascin >75 scores. High expression of PhoS39-Fascin and AcK471-Fascin was significantly associated with prolonged OS and DFS in patients with ESCC. Duncan multiple-range tests were used for Figure 1B, C, F, and G, *P < 0.05. Survival in Figure 1H was estimated by Kaplan-Meier survival analyses with the log-rank test, **P < 0.001. Abbreviations: H&E, hematoxylin and eosin; IHC, immunohistochemistry; GFP, green fluorescent protein; RFP, red fluorescent protein; WT, wild-type; FRAP, fluorescence recovery after photobleaching; ESCC, esophageal squamous cell carcinoma; LN, lymph node; SD, standard deviation; CK, cytokeratin; OS, overall survival; DFS, disease-free survival

Supplementary Figure S3). These data indicate that both Ser-39 and Lys-471 contribute to the polymerization of Factin, and phosphorylation of Fascin Ser-39 plays a more important role than the acetylation of Fascin Lys-471 in inhibiting actin-bundling activity.

To assess the clinical significance of the combination of Fascin Ser-39 phosphorylation and Fascin Lys-471 acetylation, we performed immunostaining with phosphorylation- and acetylation-specific antibodies on ESCC samples from the same cohort of patients in our previous work (Figure 1H) [6]. Importantly, Kaplan-Meier survival analysis demonstrated that high levels of both Fascin phosphorylation at the Ser-39 site and acetylation at the Lys-471 site were significantly associated with favorable overall survival (OS; P = 0.006) and disease-free survival (DFS; P = 0.003) compared with the low-level group (Figure 1H).

Here, both in vitro and in vivo biological assays showed that the cellular functions of Fascin are regulated by Lys-471 acetylation in cooperation with Ser-39 phosphorylation, suggesting a functional interplay between these two post-translational modifications. How Ser-39 phosphorylation and Lys-471 acetylation interact requires further exploration. Simultaneously targeting the modifying enzymes that regulate Ser-39 phosphorylation and Lys-471 acetylation could be a good strategy for treating esophageal cancer metastasis. Recently, we identified a novel site, Fascin Thr-403, that could regulate ESCC behavior and actin-bundling activity of Fascin through phosphorylation by AKT serine/threonine kinase 2 [8], demonstrating coordinated regulation of actin-bundling activity among Fascin's series of posttranslational modifications in order to rapidly respond to both internal and external cues. Moreover, Fascin likely regulates nuclear actin and nucleolar morphology by binding to the linker of the nucleoskeleton and cytoskeleton complex [9], and also mediates the remodeling of mitochondrial actin filaments to control mitochondrial oxidative phosphorylation [10]. Further research is needed to investigate the role of Fascin's posttranslational modifications in the nucleus and mitochondria.

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COMPETING INTERESTS

The authors declare that they have no competing interests.

AUTHORS' CONTRIBUTIONS

Fa-Min Zeng, Yin-Wei Cheng, and Jian-Zhong He designed the study. Fa-Min Zeng, Yin-Wei Cheng, Jian-Zhong He, and Xiu-E Xu conducted the experiments and analyzed the data. Fa-Min Zeng, Yin-Wei Cheng, Jian-Zhong He, Lian-Di Liao, and Xiu-E Xu acquired and analyzed the data. Fa-Min Zeng and Yin-Wei Cheng wrote the manuscript. En-Min Li and Li-Yan Xu obtained the funding to support the project and supervised the development of this work. All authors read and approved the final manuscript.

AVAILABILITY OF DATA AND MATERIALS

The data that support the findings of this study are available from the corresponding author upon reasonable request.

ETHICS APPROVAL AND CONSENT TO PARTICIPATE

This work was approved by the ethical committee of the Central Hospital of Shantou City and the ethical committee of the Medical College of Shantou University, and only resected samples from surgical patients providing written informed consent were included in this study. All animal experiments were conducted with the approval of the Institutional Animal Care and Use Committee of Shantou University.

CONSENT FOR PUBLICATION

Not applicable.

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REFERENCES

 Liu H, Zhang Y, Li L, Cao J, Guo Y, Wu Y, et al. Fascin actin-bundling protein 1 in human cancer: promising biomarker or therapeutic target? Mol Ther Oncolytics. 2021;20:240– 64.

- Lin S, Taylor MD, Singh PK, Yang S. How does fascin promote cancer metastasis? FEBS J. 2021;288(5):1434–46.
- Hashimoto Y, Kim DJ, Adams JC. The roles of fascins in health and disease. J Pathol. 2011;224(3):289–300.
- Vignjevic D, Kojima S, Aratyn Y, Danciu O, Svitkina T, Borisy GG. Role of fascin in filopodial protrusion. J Cell Biol. 2006;174(6):863–75.
- Jansen S, Collins A, Yang C, Rebowski G, Svitkina T, Dominguez R. Mechanism of actin filament bundling by fascin. J Biol Chem. 2011;286(34):30087–96.
- Cheng YW, Zeng FM, Li DJ, Wang SH, He JZ, Guo ZC, et al. P300/CBP-associated factor (PCAF)-mediated acetylation of Fascin at lysine 471 inhibits its actin-bundling activity and tumor metastasis in esophageal cancer. Cancer Commun (Lond). 2021;41(12):1398–416.
- Phair RD, Misteli T. Kinetic modelling approaches to in vivo imaging. Nat Rev Mol Cell Biol. 2001;2(12):898–907.
- Zhang ZD, Wen B, Li DJ, Deng DX, Wu XD, Cheng YW, et al. AKT serine/threonine kinase 2-mediated phosphorylation of fascin threonine 403 regulates esophageal cancer progression. Int J Biochem Cell Biol. 2022;145:106188.
- Jayo A, Malboubi M, Antoku S, Chang W, Ortiz-Zapater E, Groen C, et al. Fascin Regulates Nuclear Movement and Deformation in Migrating Cells. Dev Cell. 2016;38(4):371-83.
- Lin S, Huang C, Gunda V, Sun J, Chellappan SP, Li Z, et al. Fascin Controls Metastatic Colonization and Mitochondrial Oxidative Phosphorylation by Remodeling Mitochondrial Actin Filaments. Cell Rep. 2019;28(11):2824–36.

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