Abstract. Trefoil factor 3 (TFF3) reconstructs the epithelial barrier by stimulating epithelial cell migration and proliferation, and significantly contributes to intestinal mucosal damage and healing. In a previous study, TFF3 was identified as a novel target of microRNA-7-5p (miR-7-5p). The aim of the present study was to investigate the roles and mechanisms of miR-7-5p in the proliferation and migration of intestinal epithelial cells. Reverse transcription-quantitative polymerase chain reaction (RT-qPCR) was performed to determine the expression level of miR-7-5p in the experimental groups. In addition, western blot analysis was performed to examine the expression levels of TFF3, phosphoinositide 3-kinase (PI3K), Akt and phosphorylated (p)-AKT when miR-7-5p or TFF3 was overexpressed, and the effects of miR-7-5p and TFF3 on LS174T cell proliferation and migration were simultaneously investigated. miR-7-5p was demonstrated to decrease the expression level of TFF3, and inhibit LS174T cell proliferation and migration, which was accompanied by decreased expression levels of PI3K and p-Akt. miR-7-5p was decreased following combined treatment with the TFF3 plasmid and miR-7-5p mimics, compared with treatment with miR-7-5p mimics alone, which was accompanied by increased expression levels of TFF3, PI3K and p-Akt, and enhanced LS174T cell proliferation and migration effects. The expression levels of miR-7-5p in the miRNA negative control (NC) + LY294002 group, the miR-7-5p mimic + LY294002 group, and the miR-7-5p mimic + TFF3 plasmid + LY294002 group were higher than those in the NC group, the miR-7-5p mimic group and the miR-7-5p mimic + TFF3 plasmid group, respectively. Accordingly, the expression level of TFF3 was downregulated and the proliferation and migration ability of the cells was downregulated. The present study demonstrates that overexpressed miR-7-5p may inhibit the proliferation and migration of LS174T cells by targeting the expression of TFF3 via inhibiting the PI3K/Akt signalling pathway. The PI3K/Akt signalling pathway may exert a feedback regulation effect on miR-7-5p, inhibiting the activity of this signalling pathway, which increases the miR-7-5p expression levels and further enhances the effects of miR-7-5p on cell proliferation and migration.

Introduction

Inflammatory bowel disease (IBD), including Crohn's disease and ulcerative colitis, is characterized by chronic intestinal inflammation resulting from a complex interaction of genetically susceptible hosts, disruption of mucosal barriers, an abnormal immune response to environmental factors, and disturbance of the intestinal flora (1). The pathogenesis of IBD remains unclear, although studies in IBD animal models and IBD patients have identified a common pathologic outcome: structural and functional impairment of the intestinal mucosal barrier (2-4). The changes in intestinal epithelial mucosal barrier function are important in the occurrence, development and prognosis of IBD. Dysfunction of intestinal epithelial tight junctions leads to immune activation and promotes the occurrence of colitis (5). Clinically, the degree of mucosal repair has become the standard for evaluating the therapeutic effect of IBD (2).

The key phenomenon in the early repair of epithelial cells following injury is cell migration, whereby normal epithelial cells surrounding the damaged area migrate to the injured area, and subsequently reconstruct and maintain epithelial integrity, which is referred to as the rapid repair pathway (6). Slow repair predominantly refers to the process of cell proliferation. Cell proliferation primarily repairs the damaged mucosa by increasing the number of mitotic cells (7). Trefoil factor 3 (TFF3) is a small molecule polypeptide that is secreted by intestinal goblet cells and significantly contributes to the protection of intestinal mucosal integrity, the reconstruction and repair of intestinal mucosal injury and the anti-apoptosis of intestinal mucous cells (8,9).

Numerous studies have indicated that TFF3 promotes the migration of various types of cells in vitro and in vivo, and that the migration mechanism may be closely associated with the APC protein, β-catenin, E-cadherin, the epidermal growth factor (EGF) receptor complex and extracellular...
signal-regulated kinase (ERK). It has been found that repair of the epithelium by TFF3 is independent of transforming growth factor-β and that the epithelium is precisely repaired by promoting the migration of cells surrounding the injured area to the damaged site (10). In addition, TFF3 promotes the migration of epithelial cells by affecting the expression and localization of catenin in epithelial cells, and by inducing phosphorylation of catenin, leading to a decrease in tight junctions between cells (11,12). The EGF and Ras signalling pathways may also be involved in the mechanism through which TFF3 promotes intestinal epithelial cell migration. However, the upstream regulatory mechanism of TFF3 remains unclear.

In a previous study, TFF3 was demonstrated to be a novel target of miR-7-5p (13). miR-7 has been investigated primarily in tumour diseases. Numerous reports have indicated that miR-7 affects the functions of cell metabolism, growth, proliferation and apoptosis through acting on different target proteins (14,15). Therefore, we were interested in further examining the effects of miR-7-5p on TFF3 and the relationship among miR-7-5p, TFF3 and the proliferation and migration of intestinal epithelial cells. In the present study, we hope to provide a theoretical basis for the mechanism and treatment of IBD.

Materials and methods

Cell culture experiments. The human colonic epithelial cell line LS174T was obtained from the American Type Culture Collection (ATCC; Manassas, VA, USA). The cell line was grown to near confluence in Dulbecco’s modified Eagle’s medium supplemented with 10% fetal bovine serum (FBS; all from HyClone Laboratories, GE Healthcare Life Sciences, Logan, UT, USA), 50 µM penicillin and 50 µg/ml streptomycin at 37°C. The cells were subcultured following partial digestion with 0.25% trypsin and 0.9 mmol/l EDTA in Ca2+- and Mg2+-free phosphate-buffered saline (PBS).

TaqMan assays. To evaluate the expression level of miR-7-5p in the LS174T cells, total RNA was isolated from LS174T cells using the mirVana miRNA isolation kit (Thermo Fisher Scientific, Inc., Waltham, MA, USA). Reverse transcription (RT) was performed using the TaqMan miRNA reverse transcription kit (Thermo Fisher Scientific, Inc.) with a stem loop-specific RT primer, followed by TaqMan PCR analysis with small RNA-specific primers for miR-7-5p and U6 small nuclear RNA (snRNA; Thermo Fisher Scientific, Inc.). Briefly, RT was performed as follows: the RT Master Mix included 100 nM dNTPs with 0.15 µl dTTP, MultiScribeTM Reverse Transcription (50 U/µl; 1 µl), 10X reverse buffer (1.5 µl), 0.19 µl RNase inhibitor (20 U/µl) and 4.16 µl nuclease-free water. RT Master Mix (7 µl) was combined with 5 µl total RNA (1-10 ng), and 3 µl RT primer (5X) was added, mixed and centrifuged at 1000 x g at 4°C for 5 sec. The following parameter values were used to program the thermal cycler: 16°C temperature bath for 30 min, 42°C 30 min for RT reaction, 85°C heating for 5 min to terminate the reaction, and setting the temperature at 4°C for running qPCR. According to the manufacturer’s instructions, qPCR was performed to measure the expression levels of miR-7-5p and U6 using the Universal PCR Master Mix kit (Thermo Fisher Scientific, Inc.) and specific TaqMan probes (Thermo Fisher Scientific, Inc.). The PCR protocol was as follows: 20 µl reaction volume, including 10 µl TaqMan Universal PCR Master Mix II no UNG* (Thermo Fisher Scientific, Inc.; cat. no. 4440049), 7.67 µl nuclease-free water, 1.0 µl TaqMan Small RNA assay (20X) and 1.33 µl product from the RT reaction. The thermocycling conditions were as follows: optional AmpErase UNG activity, 50°C for 2 min; enzyme activation, 95°C for 10 min and PCR was performed for 40 cycles. Denaturing was performed at 95°C for 15 sec and extension was performed at 60°C for 60 sec. The relative expression levels of miR-7-5p were normalized to the expression of U6 RNA via the 2-∆∆Ct method (16). The measurements were performed in triplicate.

Plasmid construction and transfection. The miR-7-5p mimic, miRNA negative control (NC), miR-7-5p inhibitor and miR-7-5p inhibitor NC were purchased from Guangzhou Ribobio Co., Ltd. (Guangzhou, China). The sequences were as follows: 5'-UGGAACACAUUGAUUUUGUUU-3' for miR-7-5p mimic; 5'-UUUGUAUCACAAAGUAUGC-3' for miR-7-5p mimic NC; 5'mAmCmAmAmCmAmAmAmAmUmCmAmUmCmAmAmGmUmCmAmCmAmGmUmCmAmCmAmA-3' for miR-7-5p inhibitor; and 5'-mCmAmGmGmGmCmUmUmGmGmGmGmCmAmAmGmAmAmGmAmAmGmAmAmGmAmAmGmAmAmGmAmGmAmA-3' for miR-7-5p inhibitor NC. The LS174T cells were seeded at a density of 4x10⁴ cells/well in 6-well plates at 37°C and transfected with 100 nM miR-7-5p mimic, miRNA NC, miR-7-5p inhibitor NC, and miR-7-5p inhibitor of each at a ratio of 1:2 and 2 µg TFF3 (545.36 µg/ml). The TFF3 plasmid was purchased from SinoBiological, Inc. (Beijing, China). Transfection was performed using Lipofectamine 3000 (Thermo Fisher Scientific, Inc.) according to the manufacturer’s instructions. Cells were harvested after 48 h and subjected to various assays.

Western blot analysis. Proteins from the transfected cells were harvested at 48 h post-transfection using radioimmuneprecipitation assay buffer (50 mM Tris, pH 7.4, 150 mM NaCl, 1% Triton X-100, 1% sodium deoxycholate and 0.1% SDS and inhibitors, including sodium orthovanadate, sodium fluoride, EDTA and leupeptin, as well as proteinase/phosphatase inhibitors) for 30 min on ice. Equal quantities of protein (10-20 µg) were separated on 8-15% SDS-PAGE gels and transferred to polyvinylidene fluoride membranes (EMD Millipore, Billerica, MA, USA) at 80 v, 1.5 h. The protein expression levels of phosphoinositide-3-kinase (PI3K), Akt, phosphorylated (p)-Akt, TFF3 and glyceraldehydes-3-phosphate dehydrogenase (GAPDH) were detected using specific antibodies of anti-PI3K (rabbit polyclonal antibody; dilution, 1:1,000; cat. no. WL01169; Wanleibio, Shenyang, China), anti-AKT (rabbit polyclonal antibody; dilution, 1:1,000; cat. no. WL0003b; Wanleibio), anti-p-AKT (rabbit polyclonal antibody; dilution, 1:1,000; cat. no. WLP001; Wanleibio), anti-TFF3 (rabbit monoclonal antibody; dilution, 1:1,000; cat. no. ab108599; Abcam, Cambridge, MA, USA) and anti-GAPDH (rabbit monoclonal antibody; dilution, 1:5,000; cat. no. ab9485; Abcam). After washing in Tris-buffered saline with Tween-20 (0.1%) (Sigma-Aldrich, St. Louis, MO, USA), membranes were incubated with the secondary antibody [horseradish peroxidase-labeled goat anti-rabbit immunoglobulin G (H+L); dilution, 1:2,000; cat. no. A0208;
Cell proliferation assay. The 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay was used to measure cell proliferation. At 0, 24, 48 and 72 h post-transfection, the transfection medium was replaced with 150 µl of fresh serum-free medium containing 0.5 g/l MTT in each well. Following incubation at 37˚C for 4 h, the MTT medium was removed via aspiration, and 50 µl dimethyl sulfoxide (DMSO) was added to each well. Subsequent to incubation at 37˚C for a further 10 min, the absorbance (490 nm) of each sample was measured using a plate reader (BioTek Instruments, Inc., Winooski, VT, USA). This experiment was repeated 3 times.

Cell migration assay. The migration ability of LS174T cells was evaluated using 24-well Transwell chambers (EMD Millipore, Boston, MA, USA). For all groups, LS174T cells were diluted in serum-free medium at the logarithmic growth phase. The concentration of the cell suspension was adjusted to 5x10^4 cells/ml and 200 µl of the suspension was then inoculated into the upper Transwell chamber. The cells were subsequently incubated for 24 h at 37˚C and 5% CO₂ incubator, fixed with 4% paraformaldehyde at room temperature for 20 min and stained with 0.5% crystal violet dye for 5 min at room temperature. After washing various times with water, the number of cells that had migrated through the filter into the lower wells was counted using an inverted microscope, and 5 fields were selected and calculated the arithmetic mean value.

Experimental grouping. To illustrate the interaction of miR-7-5p, TFF3 and the PI3K/Akt signalling pathway in the regulation of LS174T cell proliferation and migration by miR-7-5p, a PI3K/Akt signalling pathway inhibitor (LY294002) and a TFF3-overexpressing plasmid were used. The LS174T cells were divided into six groups as follows: i) miRNA NC group, when the LS174T cells were 60-80% confluent, they were transfected with the miRNA mimic NC for 48 h and harvested; ii) miR-7-5p mimic experimental group, when the LS174T cells were 60-80% confluent, the cells were harvested after transfection with miR-7-5p mimics for 48 h; iii) miR-7-5p mimic + TFF3 overexpression vector transfection group, when the LS174T cells were 60-80% confluent, the cells were transfected with the TFF3 plasmid together with miR-7-5p mimics and were harvested following co-culture for 48 h; iv) miRNA mimic NC + LY294002 treatment group, 24 h after the cells were transfected with the miRNA mimic NC, they were treated with the PI3K inhibitor, LY294002 (50 µM) for 24 h; v) miR-7-5p mimic + LY294002 treatment group, 24 h after the cells were transfected with miR-7-5p mimics, they were treated with 50 µM LY294002 for 24 h; and vi) miR-7-5p mimics + TFF3 overexpression plasmid + LY294002 treatment group, the cells were transfected with miR-7-5p mimics and TFF3 plasmids for 24 h and treated with 50 µM LY294002 for 24 h.

Results

Effect of miR-7-5p on proliferation and migration of LS174T cells. The proliferation and migration functions of intestinal epithelial cells are significant following intestinal damage. The effect of miR-7-5p in the proliferation of LS174T cells was evaluated using the MTT assay, and the effect of miR-7-5p on the migration of LS174T cells was evaluated in Transwell chambers. Fig. 1 demonstrates that successful results were obtained 24 h after transfecting miR-7-5p mimics and a miR-7-5p inhibitor. As presented in Fig. 2, compared with the NC group, the miRNA mimic NC group and the miRNA
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Effect of miR-7-5p on the expression levels of TFF3, PI3K, Akt and p-Akt protein. As shown in Fig. 4, compared with the NC group, the miRNA mimic NC group and the miRNA inhibitor NC group, the overexpression of miR-7-5p decreased the levels of TFF3, PI3K and p-Akt protein expression, whereas inhibiting the expression level of miR-7-5p increased TFF3, PI3K and p-Akt protein expression levels and the difference was statistically significant (P<0.05). However, no effect of miR-7-5p was observed on the expression level of the Akt protein. In a previous study, miR-7-5p was demonstrated to bind to the 3'UTR of TFF3. Thus, these results indicated that miR-7-5p influences the proliferation and migration of LS174T cells by targeting TFF3, with the possible participation of the PI3K/Akt signaling pathway.

Effect of miR-7-5p, TFF3 and LY294002 on the protein expression levels of TFF3, PI3K, Akt and p-Akt. According to
the experimental groups, the transfected cells were harvested 48 h after transfection, and the mRNA expression level of miR-7 was detected via qPCR. As demonstrated in Fig. 5, the expression level of miR-7-5p in the miR-7-5p mimic group was significantly higher than that in the miRNA NC group. Compared with the miR-7-5p mimic group, the expression level of miR-7-5p in the miR-7-5p mimic + TFF3 plasmid group was significantly decreased, while compared with the miR-7-5p mimic + TFF3 plasmid group, the expression level of miR-7-5p in the miR-7-5p mimic + LY294002 group was significantly increased, and this difference was statistically significant (P<0.05). These findings indicated that the TFF3 overexpression plasmid was able to inhibit the expression level of miR-7-5p, but that LY294002 inhibited this inhibitory effect of the TFF3 plasmid on miR-7-5p. The expression level of miR-7-5p in the miRNA NC + LY294002 group, the miR-7-5p mimic + LY294002 group, and the miR-7-5p mimic + TFF3 plasmid + LY294002 group was greater than that in the NC group, the miR-7-5p mimic group, and the miR-7-5p mimic + TFF3 plasmid group, respectively, and these differences were statistically significant (P<0.05). Thus, these findings indicated that the expression level of miR-7-5p was upregulated by the PI3K/Akt signalling pathway inhibitor, indicating that inhibition of the PI3K/Akt signalling pathway may result in feedback upregulation of miR-7-5p expression.

The TFF3 protein was detected by western blot analysis (Fig. 6). The results (Figs. 6-8) showed that the expression levels of the TFF3, PI3K, and p-Akt proteins in the miR-7-5p mimic group were significantly lower than those in the NC group, whereas the expression levels of the TFF3, PI3K and p-Akt proteins in the miR-7-5p mimic + TFF3 plasmid group were higher than those in the miR-7-5p mimic group. The difference between these groups was significant (P<0.05), indicating that miR-7-5p downregulates the protein expression of its target, TFF3. The TFF3 promotes PI3K protein expression, and miR-7-5p inhibits the activation of the PI3K/Akt signalling pathway while downregulating TFF3 protein expression levels. However, following treatment of the miR-7-5p mimic + TFF3 plasmid group with LY294002, the expression levels of the TFF3, PI3K and p-Akt proteins decreased, indicating that blocking the PI3K/Akt signalling pathway further promotes the effect of miR-7-5p on TFF3. Compared with the NC group, the expression level of the TFF3 protein in the NC + LY294002 group was significantly decreased (P<0.05). Compared with the miR-7-5p mimic group, the expression level of the TFF3 protein in the miR-7-5p mimic + LY294002 group was significantly decreased (P<0.05). Compared with the miR-7-5p mimic + TFF3 plasmid group, the expression level of the TFF3 protein in the miR-7-5p mimic + TFF3 plasmid + LY294002 group was also significantly decreased (P<0.05). Taken together, these findings demonstrated that subsequent to adding the PI3K/Akt signalling pathway inhibitor, the expression level of miR-7-5p was upregulated and the expression level of TFF3 was downregulated, which
indicated that inhibiting the PI3K/Akt signalling pathway may exert a feedback effect leading to the upregulation of miR-7-5p expression and the inhibition of TFF3 expression.

The expression levels of the PI3K and p-Akt protein were consistent with the trend of TFF3 protein expression (Figs. 7 and 8), although there was no significant difference in the expression levels of the AKT protein between the groups (Fig. 8B), suggesting that the primary mechanism by which miR-7-5p regulates the expression level of TFF3 and affects the PI3K/Akt signalling pathway occurs via modulation of Akt protein phosphorylation.

Effect of miR-7-5p, TFF3 and LY294002 on the proliferation and migration of LS174T cells. To detect the cell proliferation of each group, the cell optical density (OD) at 490 nm was detected at 0, 24, 48 and 72 h after transfection. As shown in Fig. 9, compared with the NC group, the OD value of the miR-7 mimic group decreased significantly, indicating a decreased cell proliferation ability; the difference between the groups was statistically significant (P<0.05). Compared with the miR-7-5p mimic group, the miR-7-5p mimic + TFF3 plasmid group showed a significantly increased OD value and enhanced cell proliferation ability, with a significant difference.
observed between the groups (P<0.05). Compared with the miR-7-5p mimic group, the miR-7-5p mimic + LY294002 group exhibited a slight decrease in the OD value, although the difference between the groups was not significant (P>0.05). This result indicates that in addition to the PI3K/Akt signalling pathway, miR-7-5p affects cell proliferation via pathways other than regulating TFF3. However, compared with the miR-7-5p mimic + TFF3 plasmid group, the miR-7-5p mimic + TFF3 plasmid + LY294002 group exhibited a decreased OD value and decreased cell proliferation ability. These results indicated that the PI3K/Akt signalling pathway is involved in the regulation of TFF3 by miR-7-5p, subsequently affecting the proliferation of LS174T cells, and that blocking the PI3K/Akt signalling pathway further promotes the inhibitory effect of miR-7-5p on cell proliferation.

The present study illustrated the effect of miR-7-5p, TFF3 and LY294002 on the migration of LS174T cells via Transwell chamber assays. The number of cells that migrated through the Transwell chamber was counted following 24 h of incubation. As presented in Fig. 10, compared with the NC group, the cell migration ability of the miR-7-5p high expression group was significantly decreased. Compared with the miR-7-5p mimic group, the number of cells passing through the Transwell membrane was increased in the miR-7-5p mimic + TFF3 plasmid group. This finding indicated that the cell migration ability was increased, indicating that the TFF3 plasmid weakened the inhibitory effect of miR-7-5p on the migration ability of the cells, and supported the effect of miR-7-5p by targeting TFF3 and mediating cell migration. Compared with the miR-7-5p mimic group, the miR-7-5p mimic + LY294002 group exhibited a decrease in the number of cells passing through the Transwell membrane, suggesting that LY294002 further downregulated the migration ability of LS174T cells. The number of cells passing through the Transwell membrane in the miR-7-5p mimic group decreased, whereas it increased in the miR-7-5p mimic + TFF3 plasmid group. However, when the miR-7-5p mimic + TFF3 overexpression plasmid group was treated with LY294002, the number of cells passing through the Transwell membrane decreased once more, indicating that LY294002 blocked the promoting effect of the TFF3 overexpression plasmid on cell migration. Taken together, these findings indicate that the effect of miR-7-5p on the migration of LS174T cells via TFF3 regulation was mediated by the PI3K/Akt signalling pathway, and blocking this pathway may further promote the inhibitory effect of miR-7-5p on cell migration.
Discussion

miRNAs are a type of snRNA that lead to mRNA degradation and translational repression by binding to the 3'UTR of target mRNAs. miRNAs are significant in processes, such as cell proliferation, apoptosis, growth and development, cell differentiation and metabolism (17).

miR-7 is one type of mature miRNA that has been investigated in tumour diseases. Numerous reports have suggested that miR-7 affects the functions of cell metabolism, growth, proliferation and apoptosis by acting on different target proteins (14,15,18-20). Fang et al (18) proposed that miR-7 inhibits the growth and metabolism of hepatocellular carcinoma by regulating the PI3K/Akt signalling pathway. Meza-Sosa et al (19) demonstrated that miR-7 promotes epithelial cell transdifferentiation by targeting Kruppel like factor 4. Xu et al (20) revealed that miR-7 regulates XRCC2, and thereby inhibits the proliferation of colorectal cancer cells and induces apoptosis (20). In our previous study, miR-7-5p was found to be differentially expressed in IBD lesions and normal tissues. In addition, it was confirmed that miR-7-5p binds to the TFF3 3'UTR and regulates the expression level of TFF3 at the post-transcriptional level (13).

The TFF3 proteins are a group of small molecular peptides that are predominantly secreted by goblet cells (21). TFF3 exhibits a special three-lobed structure. This structure exhibits high stability, which causes the TFF to present strong anti-protease hydrolysis, acid digestibility and heat-resistant properties. Therefore, TFF3 maintains biological activity in the complex environment of the digestive tract (22). Investigation of the response of TFF3 to mucosal damage and its response in patients with IBD has indicated that TFF3 is significant in injury repair and mucosal protection, and the underlying mechanisms include the promotion of cell migration, cell proliferation and anti-apoptosis effects (23-25).

The present study demonstrates that miR-7 may regulate the function of tumour cells by regulating multiple pathways, such as the PI3K/Akt and mitogen-activated protein kinase kinase/ERK signalling pathways (26,27). Li et al (28) found that miR-7 inhibits the proliferation and invasion of human colorectal cancer cells, and that PI3K signalling pathways are involved in this process. Zhang et al (29) showed that miR-21 could regulate the tight junctional permeability of intestinal epithelial cells, which was mediated by the PI3K/Akt signalling pathway. It has also been suggested that TFF3 and the PI3K/Akt pathway play important roles in cell migration and proliferation. Dise et al (30) demonstrated that EGF promoted intestinal epithelial cell migration by activating PI3K and causing Rac activation. Langlois et al (31) showed that PTEN inhibits tumour progression by controlling cell polarity, the establishment of cell-cell junctions, paracellular permeability, migration and metabolic potential and that PTEN phosphorylation is mediated by the activation of PI3K. Sun et al (32) demonstrated that intestinal trefoil factor (ITF) can promote the proliferation and migration of gastric epithelial cells. ITF protects the integrity of gastric epithelial cells from damage by activating PI3K/Akt cell signalling pathways. Lin et al (33) revealed that TFF3 overexpression may promote the expression of zonula occludens-1, occludin and claudin-1, but this effect was inhibited after suppressing the PI3K/Akt signalling pathway.

Therefore, the aim of the present study was to elucidate the association between miR-7-5p, TFF3 and PI3K/Akt in the process, whereby miR-7-5p regulates the proliferation and migration of intestinal epithelial cell. The current results revealed that miR-7-5p decreased the expression level of TFF3, and inhibited LS174T cell proliferation and migration, accompanied by decreased expression of PI3K and p-Akt. Furthermore, it was identified that miR-7-5p expression levels were decreased following combined treatment with the TFF3 plasmid and miR-7-5p mimics, compared with treatment with the miR-7-5p mimics alone. This effect was accompanied by increased expression levels of TFF3, PI3K and p-Akt, and by enhanced LS174T cell proliferation and migration. Furthermore, the expression level of miR-7-5p in the miRNA NC + LY294002 group, the miR-7-5p mimic + LY294002 group, and the miR-7-5p mimic + TFF3 plasmid + LY294002 group was higher than in the NC group, the miR-7-5p mimic
group, and the miR-7-5p mimic + TFF3 plasmid group, respectively. Accordingly, the expression level of TFF3 was downregulated, and cell proliferation and migration were downregulated simultaneously. These data indicate that miR-7-5p regulates TFF3 and inhibits the proliferation and migration of LS174T cells. The PI3K/Akt signalling pathway is involved in the regulation of cell proliferation and migration via miR-7-5p targeting TFF3 (Fig. 11). The PI3K/Akt signalling pathway may exert a feedback regulation effect on miR-7-5p; inhibition of the activity of this pathway may enhance miR-7-5p expression levels and further enhance the effect of miR-7-5p on cell proliferation and migration.

In conclusion, miR-7-5p inhibits the proliferation and migration of LS174T cells by targeting TFF3 via inhibiting the PI3K/Akt signalling pathway. Additionally, the PI3K/Akt signalling pathway may have a feedback regulation effect on miR-7-5p. Therefore, miR-7-5p may serve as a therapeutic target for protection of intestinal epithelial barrier integrity.

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