miR-210-3p suppresses osteogenic differentiation of MC3T3-E1 by targeting brain derived neurotrophic factor (BDNF)

Li Deng¹,², Shuang Lai¹, Liyuan Fan¹, Xinlun Li³, Hao Huang⁴ and Yandong Mu¹*

Abstract

Background and objective: As an important mediator of intercellular interaction and formation of extracellular bone matrix, porous scaffolds are widely used for bone regeneration. Accumulating evidences demonstrate that microRNA are involved in the regulation of scaffolds-induced bone regeneration. Recently, we revealed that miR-210-3p was highly expressed during osteogenesis induced by HAG. In present study, we further explored the molecular mechanism underlying the effect of miR-210-3p on osteogenic differentiation.

Materials and methods: In this study, miR-210-3p mimics and inhibitors were synthesized and transfected into MC3T3-E1 cells to explore their effects on osteogenic differentiation. The expression of osteogenic marker (Alp and Runx2) were detected by real-time quantitative PCR (qRT-PCR) and western blotting. After osteogenesis induction for 7 days, Alp staining were used to detected osteoblast differentiation of MC3T3-E1 cells. CCK8 and Transwell assays were performed to detected cell proliferation and migration. Then, top ranking list of target genes of miR-210-3p obtained from TargetScan and the expression of BDNF were detected by qRT-PCR and ELISA. The relationship between miR-210-3p and BDNF was verified by luciferase report assay. Furthermore, the effect of BDNF on osteoblast differentiation was verified by transfecting siRNA or adding BDNF to the culture medium.

Results: MiR-210-3p mimics markedly suppress osteogenic differentiation, cell migration and cell proliferation of MC3T3-E; nevertheless, silencing of miR-210-3p dramatically enhanced MC3T3-E1 osteogenesis, cell migration and proliferation. Furthermore, luciferase reporter assay verified that brain derived neurotrophic factor (BDNF) is a directly target of miR-210-3p. Moreover, BDNF siRNA significantly decreased the expression levels of ALP and cell migration. The addition of BDNF partially rescued the inhibition of osteogenesis by miR-210-3p.

Conclusion: miR-210-3p inhibited the osteogenic differentiation via targeting BDNF. Our Results provide a promising target for regulating osteogenic differentiation.

Keywords: miR-210-3p, Osteogenic differentiation, Cell migration, BDNF

Introduction

Bone defects caused by different causes lower people’s quality of life and increase economic burden. Despite the regenerative potential of bone, the repair of large bone defects is still a major clinical problem. Autografts, as the gold standard for bone defect repair, has disadvantages such as limited bone graft volume, potential donor site

*Correspondence: muyd@uestc.edu.cn

¹ Stomatology Department, Sichuan Provincial People's Hospital, School of Medicine, University of Electronic Science and Technology of China, Chengdu 611731, China
Full list of author information is available at the end of the article

© The Author(s) 2022. Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article’s Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article’s Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated in a credit line to the data.
lesions and wound infection [1–3]. Therefore, bone tissue engineering is widely used in bone regeneration as the most promising method. Scaffold is an important part of bone tissue engineering. In previous study, we developed a grooved porous hydroxyapatite scaffold (HAG) with good osteogenic effect in vitro and in vivo [4].

As a mineralized mesenchymal tissue, bone regeneration is a complex process involving the balance of various cell differentiation [5]. This process is extensively regulated. MicroRNAs (miRNAs), a group of non-coding RNA consisting of 20–24 nucleotides are evolutionarily conserved, interaction of target mRNA sequences, negative regulation of their expression and participate in biological processes including cell differentiation and proliferation [6–8]. At present, they have been confirmed to be tightly involved in the regulation of bone formation by targeting different target genes and negatively regulating their expression [9]. MiR-23a, miR-30c, miR-34c and miR-133a et al. have been confirmed that significantly inhibit osteoblast differentiation by targeting RUNX2 (RUNX family transcription factor 2, an essential regulatory factor for skeletal morphogenesis) [10]. MiR-21 promotes BM-MSCs osteogenesis by targeting an TGF signal antagonist SMAD7 [11]. Upon the development of bone tissue engineering, the important role of miRNA in scaffold osteogenesis is increasingly being revealed [12–14]. Maryam Izadpanahi et al. investigated the nanorographic niche interactions with non-coding RNA in stem cell fate to reveal deeper mechanisms for enhancing osteogenic differentiation. They found that the activity of miRNA in hMSCs was closely related to the nanotopographical cues and that the nanotopographical cues regulated the osteogenic differentiation of hMSCs via modulating IncRNAs [15]. Recently, we found that miRNAs were differentially expressed during osteogenesis induced by HAG scaffolds, and miR-210-3p was one of the significantly up-regulated miRNAs [16]. However, its role of osteogenic differentiation remains unclear.

In the present study, the role and mechanism of miR-210-3p in the osteogenic differentiation were investigated. Our findings could provide new insights and target for the diagnosis and treatment of bone regeneration.

**Materials and methods**

**Cell culture and transfection**

MC3T3-E1 cell line were obtained from the Cell Bank of Typical Culture Preservation Committee of the Chinese Academy of Science, Shanghai, China. Cells were cultured in complete medium (MEM (Hyclone) with 10% fetal bovine serum (FBS, Avantor®) and 1% penicillin streptomycin (HyClone)).

MiR-210-3p mimic, inhibitor and BDNF siRNA (synthesized by GenePharma, 100 pM) were transfected when the cell density approached 80% using the Lipofectamine™ 2000 (Thermo Fisher Scientific) transfection reagent following the manufacturer’s protocol. After 24 h, the medium was replaced with osteogenic induction medium. 20 mM β-glycerophosphate and 50 g/mL vitamin C were added to complete medium for osteogenic induction. Change the medium every three days. After induction for 7 days, Alkaline phosphatase (ALP) expression was detected according to the manufacturer’s instructions (TRAP/ALP Stain Kit, Wako).

**Quantitative real-time PCR**

TRIzol™ reagent (Invitrogen) was used for total RNA extraction. M-MLV (Thermo Fisher Scientific) was used for reverse transcription of RNA. Realtime PCR were performed by ABI 7500 (with 2X ChamQ Universal SYBR Master Mix (Vazyme, China)). The primer sequences used are as follows:

| Gene | Forward (5’-3’) | Reverse (5’-3’) |
|------|----------------|----------------|
| miR-210-3p | ACTGTCGGCTGTGACAGC | GAGAGGAGAAGGAAGGGGA |
| Alp | GCAGTTATGATGAAACTGCAGAACAC | ATGGCCCTGGATCCTCACCAC |
| Gapdh | AGGTCGGGTGTGAAGC | TTGATGACCATAGTGTG |

**Protein extraction and western blot analysis**

Total protein was extracted using RIPA lysis buffer (Beyotime), and the proteins were quantitatively determined by BCA kit (Biosharp). 10% SDS-PAGE gel electrophoresis was used to separate protein samples for Western blot analysis. The primary antibodies used were anti-ALP (HuaBio, 1:2000), Runt-related transcription factor 2 (RUNX2) (HuaBio, 1:2000), GAPDH (HuaBio, 1:5000). The HRP-conjugated secondary antibody was used to enable detection. The super-ECL detection system (Biosharp) was used in combination with primary antibody.

**Enzyme-Linked Immunosorbent Assay (ELISA).**

The Mouse BDNF (Brain Derived Neurotrophic Factor) ELISA Kit (Elabscience) was used to measure the protein level of BDNF in the medium. The culture supernatants were collected and centrifuged for 20 min at 1000× g. Then processed according to the manufacturer’s instructions and measured the absorbance at 450 nm.

**Cell migration and proliferation**

Transwell chambers (8 µm pores, Corning) were used to detect cell migration. In details, the MEM cell suspension (5 × 10⁵ cells) was seeded in the chamber and the lower filled with complete medium. After 24 h and 48 h
miR-210-3p inhibited the osteogenic differentiation of MC3T3-E1 cells. A and B Expression of miR-210-3p after transfection with mimics or inhibitor were examined by qPCR. C and D The mRNA expression of Alp was detected after transfection with mimics or inhibitor and osteogenesis induction for 3 days and 6 days. E and F The protein expression of Alp and Runx2 were detected after transfection with mimics or inhibitor. G Representative images of Alp staining after transfection and Columnar analysis diagram (H)
Fig. 2  miR-210-3p inhibited the proliferation and migration of MC3T3-E1 cells. A The migration of MC3T3-E1 cells transfected with miR-210-3p mimic or inhibitor was evaluated, assessed by staining, photographed and Columnar analysis diagram (B). C The proliferation of MC3T3-E1 cells transfected with miR-210-3p mimic or inhibitor was detected by CCK-8 assay and cell images at day 5 (D)
of culture, the MEM in the chamber was discarded, and cotton wiped cells from the upper chamber. The bottom cells were stained with crystal violet after fixation with 4% PFA for 15 min. Finally, the cells were photographed and analyzed by image J. For proliferation detected, cells were seeded in 96-well plate after transfection. According to manufacturer’s protocol, a CCK8 assay (Solarbio) was performed on day 3 and 5. An absorbance of 450 nm represents the cell proliferation ability. The experiments were repeated three times.

Dual luciferase reporter assay
Target scan predicted the 3' untranslated region (UTR) binding site of BDNF to miR-210-3p. Luciferase assays were performed to verify the interaction between miR-210-3p and BDNF. Briefly, BDNF luciferase reporter plasmid (constructed by GenePharma) was transfected into MC3T3-E1 cells with miR-NC or miR-210-3p mimic. The medium was replaced after 4 h. After incubated for 24 h, a Dual-Luciferase Reporter Test Kit (Vazyme) was used to measure the Firefly and Renilla luciferase activities. The ratio of firefly to Renilla luciferase activity is the relative luciferase activity of each sample. The experiments were repeated three times.

Statistical analysis
GraphPad Prism 8 was used for statistical analyses. Student’s t test was conducted to compare means between two groups. All data are expressed as the mean ± standard deviation. P values < 0.05 were considered statistically significant.

Results
miR-210-3p inhibited the osteogenic differentiation of MC3T3-E1 cells
Our previous study revealed that miR-210-3p was upregulated in HAG scaffold induced osteogenic differentiation [16]. miR-210-3p is a highly conserved miRNA from mouse to humans (sequence: CUGUCGUGUGACAG CGGCUGA). For uncovering the role of miR-210-3p in osteogenic differentiation, we first achieved efficient overexpression or silencing of miR-210-3p in cells by transfection, as shown in Fig. 1A and B. Then, we further detected osteogenic markers expression. Surprisingly, the results showed that miR-210-3p overexpression decreased Alp mRNA expression, while miR-210-3p inhibitors increased it (Fig. 1C and D). Consistently, the protein expression of Alp and Runx2 were decreased or increased with miR-210-3p mimic or inhibitor transfection (Fig. 1E and F). Moreover, ALP staining also showed that miR-210-3p mimic significantly decreased ALP expression while inhibitor promoted (Fig. 1G and H). These results suggested that miR-210-3p inhibited the osteogenic differentiation of MC3T3-E1 cells.

miR-210-3p inhibited the proliferation and migration of MC3T3-E1 cells
Cell proliferation and migration have been proved to be closely related to their osteogenic differentiation ability. Here, cell migration of MC3T3-E1 was detected after transfection with miR-210-3p mimic or inhibitor. As shown in Fig. 2A and B, crystal violet staining showed that miR-210-3p overexpression inhibited the migration of MC3T3-E1 cells cultured 24 h or 48 h in Transwell chambers, while inhibitors promoted. Similarly, the results of CCK8 assay showed that highly expression of miR-210-3p reduced cell proliferation of MC3T3-E1 cells (Fig. 2C). Cell images were photoed on the fifth day after culture and showed that miR-210-3p significantly negatively regulated cell proliferation (Fig. 2D).

BDNF is a directly target of miR-210-3p
To uncover the downstream molecular mechanism contributing to miR-210-3p mediated osteogenesis, TargetScan software was used to identify potential targets of miR-210-3p. Notably, we found that BDNF as an important target of miR-210-3p (Fig. 3A). Furthermore, we detected the mRNA and protein expression of Bdnf after miR-210-3p transfection to validate the prediction. Indeed, as shown in Fig. 3B and C, Bdnf decreased after miR-210-3p overexpression, while increased after silencing. Moreover, we constructed the BDNF luciferase reporter plasmid, which contained either wild type fragments (BDNF-wt) or mutant binding site fragments (BDNF-mut) downstream of the firefly sequence (Fig. 3D). The miR-210-3p mimic was co-transfected into MC3T3-E1 cells with the constructed reporter plasmids.

(See figure on next page.)

Fig. 3 BDNF is a directly target of miR-210-3p. A Top ranking list of miR-210-3p target genes obtained from TargetScan software. B The mRNA expression of Bdnf after transfection with miR-210-3p mimics were examined by qRCR. C Relative protein expression of Bdnf were examined after transfection with miR-210-3p mimics or inhibitor. D Binding sites of miR-210-3p and the BDNF 3'UTR, as detected by luciferase reporter assays. BDNF mutation: luciferase reporter plasmid containing mutant BDNF 3'UTR. BDNF-WT: luciferase reporter plasmid containing wild-type BDNF 3'UTR. E Relative luciferase activities of luciferase reporters containing WT or MUT BDNF 3'UTR in MC3T3-E1 cells transfected with miR-210-3p mimic.
**Fig. 3** (See legend on previous page.)

### A. Top ranking list target gene (Target scan)

| Target gene | Gene name | 3P-seq tags + 5 | Aggregate PCT |
|-------------|-----------|----------------|---------------|
| C6orf136    | chromosome 6 open reading frame 136 | 307            | 0.17          |
| GPD1L       | glyceraldehyde-3-phosphate dehydrogenase 1-like | 130 | 0.61          |
| B4GALT5     | UDP-Gal:betaGlcNAc beta 1,4-
| galactosyltransferase, polypeptide 5 | 1000 | 0.17          |
| ISCU        | iron-sulfur cluster assembly enzyme | 182            | 0.27          |
| LIFN2       | extracellular leucine-rich repeat and fibronectin type III domain containing 2 | 22 | 0.47          |
| KLF7        | Kruppel-like factor 7 (ubiquitous) | 737            | 0.47          |
| NDUFA4      | NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4.9 kDa | 143 | 0.21          |
| CYGB        | cytoglobin | 10             | 0.27          |
| SYNGAP1     | synaptic Ras GTPase activating protein 1 | 15 | 0.17          |
| BDNF        | brain-derived neurotrophic factor | 2696         | 0.21          |

### B. 

- **View 1**: Bar graph showing the mRNA expression of Bdnf in NC, miR-210-3p mimics, and miR-210-3p mimics.
- **View 2**: Bar graph showing the relative protein expression of BDNF in NC, mimics, and inhibitors.

### D. 

- **View 1**: Diagram of Mouse Bdnf ENSG00000017669 3' UTR length: 2985.
- **View 2**: Diagram showing the BDNF-WT and BDNF-mutation with the miR-210-3p binding site.

### E. 

- **View 1**: Bar graph showing the relative luciferase activity (%) for NC, BDNF-WT, BDNF-mutation, BDNF-WT, and BDNF-mutation.

(See legend on previous page.)
BDNF promoted osteoblast differentiation, cell migration and partially rescued the miR-210-3p mediated inhibition of osteogenesis. A and B Western blot analysis of protein expression of Alp (A. BDNF added; B. BDNF rescue). C Relative protein expression of BDNF and Alp were examined after transfection with BDNF siRNA. D and F. The images of cell migration and Alp staining (D. BDNF added; E. BDNF rescue; F. BDNF siRNA transfection)
The results showed that compared to BDNF-mut, co-transfection of BDNF-wt with miR-210-3p mimic decreased luciferase activity (Fig. 3E). These data imply that BDNF is a directly target of miR-210-3p.

**BDNF promoted osteoblast differentiation, cell migration and partially rescued the miR-210-3p mediated inhibition of osteogenesis**

In order to further reveal the role of miR-210-3p and BDNF in osteogenic differentiation of MC3T3-E1 cells, we first added different concentrations of BDNF in the medium to induce osteogenic differentiation. Western blotting showed that BDNF-added increased the expression of ALP (Fig. 4A), and the BDNF-added could partially rescue inhibitory effect of miR-210-3p on ALP expression (Fig. 4B). Furthermore, transfection of BDNF siRNA silenced the expression of BDNF and decreased the expression of Alp (Fig. 4C). The promoting effect of BDNF on MC3T3-E1 cell proliferation and ALP expression was also confirmed by ALP staining and cell migration assay ALP staining and transwell assay also showed that BDNF could promote the osteogenic differentiation and cell migration of MC3T3-E1 cells (Fig. 4D). Meanwhile, the BDNF-added also could partially rescue the inhibitory effect of miR-210-3p on ALP expression and cell migration in MC3T3-E1 (Fig. 4E). Consistently, the silencing of BDNF inhibits ALP expression and cell migration of MC3T3-E1 cells (Fig. 4F). These data confirmed that miR-210-3p inhibited the osteogenic differentiation of MC3T3-E1 cells by targeting BDNF.

**Discussion**

The osteogenic differentiation ability of cells is essential for bone regeneration. Increasing evidences suggest that miRNAs are important and functional in regulating osteogenesis [17, 18]. Hassan et al. and Zhang et al. found that miR-218 has positively effects on the Wnt signaling pathway, through targeted the signaling different inhibitors, including DKK2, SFRP2 and SOST [19, 20]. Meanwhile, SOST also was targeted and inhibited by miR-96 [21]. MiR-146a was found to negatively regulate osteogenesis and bone regeneration both in vitro and in vivo through interaction with SMAD4 [22]. However, the role of miR-210-3p in osteogenic differentiation remains unclear. In current study, our data exhibit that miR-210-3p obviously inhibited the osteogenic differentiation and cell migration of MC3T3-E1 cells by targeting BDNF.

After bone graft, the loss of neuralized vascular network in the implanting area would affect subsequent bone regeneration. Brain-derived neurotrophic factor (BDNF) is a widely recognized growth factor involved in the regulation of neuronal growth, survival and angiogenesis [23]. Moreover, BDNF seems to be considered a growth factor suitable for bone material implantation. Yamashiro et al. and Kilian et al. showed that the mRNA and protein of BDNF existed in osteoblasts and were increased during human fracture healing [24]. In addition, the role of BDNF in promoting osteogenic differentiation has been confirmed both in vitro and vivo [25, 26]. Consistently, in our study, we confirmed that BDNF promoted the osteogenic differentiation and cell migration of MC3T3-E1. Furthermore, the addition of BDNF can partially rescue the inhibition caused by miR-210-3p.

Recently, miRNA has also been shown to play a vital regulatory role in tendon healing and osteoarthritis (OA), involving apoptosis, senescence of tendon stem/progenitor cell, and maintenance of chondrocyte homeostasis [27, 28]. Furthermore, based on the natural processes of regulation of eukaryotic genes. Small interfering RNAs (siRNAs) mediated gene silencing has been used to regulate tendon homeostasis and human rheumatoid arthritis [29, 30]. Consistently, in present study, the functional role of miR-210-3p may introduce the idea of using miR-210-3p targeted silencing-modified cell or scaffolds for bone repair. Meanwhile, the expression of miR-210-3p may be a prognostic indicator of bone regeneration. Moreover, BDNF transfer may be an efficient way to bone regeneration through regulate bone metabolism and promote osteogenesis.

In conclusion, our study has shown a clear link between miR-210-3p and BDNF in MC3T3-E1 cells and demonstrated that miR-210-3p inhibited osteogenic differentiation by targeting BDNF. This suggests that silencing miR-210-3p or overexpressing BDNF may have a future role in improving bone tissue engineering.

**Authors contribution**

Li Deng. wrote the main manuscript text and prepared Figs. 3–4. Yandong Mu. provide Funding support and design experiments, Shuang Lai. and Liyuan Fan. prepared Figs. 1, Xinlun Li and Hao Huang prepared Figs. 2. All authors reviewed the manuscript.

**Funding**

This work was supported by National Natural Science Foundation of China (82071168), the Science and Technology Department of Sichuan Province Foundation (2021YFS0009).

**Availability of data and materials**

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

**Declarations**

**Ethics Approval and consent to participate**

Not applicable.

**Competing interests**

The authors declare that they have no competing interests.
Author details
1 Stomatology Department, Sichuan Provincial People’s Hospital, School of Medicine, University of Electronic Science and Technology of China, Chengdu 611731, China. 2 Institute of Tissue Engineering and Stem Cells, Nanchong Central Hospital, The Second Clinical College of North Sichuan Medical College, Nanchong 637000, Sichuan, China. 3 Stomatology Department, Sichuan Provincial People’s Hospital, Chengdu 610072, Sichuan, China. 4 Key Laboratory of Advanced Technologies of Materials (Ministry of Education), School of Materials Science and Engineering, Southwest Jiaotong University, Chengdu 610031, Sichuan, China.

Received: 11 July 2022. Accepted: 31 August 2022
Published online: 14 September 2022

References
1. Keating J, McQueen M. Substitutes for autologous bone graft in orthopaedic trauma. J Bone Joint Surg Br Vol. 2001;83(1):3–8.
2. Finkemeier CG. Bone-grafting and bone-graft substitutes. J Bone Joint Surg Am. 2002;84(4):544–64.
3. Giannoudis PV, Einhorn TA, Marsh D. Fracture healing: the diamond concept. Injury. 2007;38:53–6.
4. Ren X, Qiang T, Tian K, et al. Enhancement of osteogenesis using a novel porous hydroxyapatite scaffold in vivo and vitro. Ceram Int. 2018;44:21656–65.
5. Kapinas K, Delany AM. MicroRNA biogenesis and regulation of bone remodeling. Arthritis Res Ther. 2011;13:220.
6. Goff LA, et al. Differentiating human multipotent mesenchymal stromal cells regulate microRNAs: prediction of microRNA regulation by PDGF during osteo-genesis. Exp Hematol. 2008;36(10):1354-1369 e2.
7. Ribeiro OA, et al. MicroRNAs: modulators of cell identity, and their applications in tissue engineering. MicroRNA. 2014;3(1):45–53.
8. Vimalraj S, Selvamurugan N. MicroRNAs expression and their regulatory networks during mesenchymal stem cells differentiation toward osteoblasts. Int J Biol Macromol. 2017;96:144–202.
9. Lian JB, Stein GS, van Wijnen AJ, Stein JL, Hassan MQ, Gaur T, Zhang Y. MicroRNA control of bone formation and homeostasis. Nat Rev Endocrin. 2012(8):212–27.
10. Zhang Y, Xie R-J, Croce CM, et al. A program of microRNAs controls osteogenic lineage progression by targeting transcription factor Runx2. Proc Natl Acad Sci. 2011;108:9863–8.
11. Li X, Guo L, Liu Y, et al. MicroRNA-21 promotes osteogenesis of bone marrow mesenchymal stem cells via the Smad7-Smad1/5/8-Runx2 pathway. Biochem Biophys Res Commun. 2017;493:928–33.
12. Pizzicannella J, Cavalcanti M, Trubian O, Diomede F. MicroRNA 210 mediates osteogenic differentiation and bone formation in ankylosing spondylitis mice through activating the Wnt signaling pathway by binding to SOST. J Cell Biochem. 2019;120:15429–42.
13. Xie Q, Wei W, Ruan J, Ding Y, Zhaoua A, Bi X, Sun H, Gu P, Wang Z, Fan X. Effect of miR-146a on the osteogenesis of adipose-derived mesenchymal stem cells and bone regeneration. Sci Rep. 2017;7:42840.
14. Kermani P, Hempstead B. Brain-derived neurotrophic factor: a newly described mediator of angiogenesis. Trends Cardiovasc Med. 2007;17:140–3.
15. Yamashiro T, Fukunaga T, Yamashita K, Kobashi N, Takano-Yamamoto T. Gene and protein expression of brain-derived neurotrophic factor and TrkB in bone and cartilage. Bone. 2001;28:404–9.
16. Kaucshik V, Gebert A, Calin M, Eckert J, Scheich S, Heiss C, Lips KS. Effects of new beta-type Ti-60N implant materials, brain-derived neurotrophic factor, acetylcysteine and nicotine on human mesenchymal stem cells of osteoprotic and non osteoporotic donors. PLoS ONE. 2018;13: e0193468.
17. Kaucshik V, Schneider M, Jauch A, Schumacher M, Kampschulte M, Rohnke M, Herss A, Bambegg C, Trinkle K, Gelinsky M, et al. Effects of a partly bone cement containing brain-derived neurotrophic factor-functionalized mesoporous bioactive glass particles on metaphysical healing in a new murine osteoporotic fracture model. Int J Mol Sci. 2018;19:3531.
18. Giordano L, Porta GD, Peretti GM, Maffulli N. Therapeutic potential of microRNA in tendon injuries. Br Med Bull. 2020;103(1):79–94.
19. Oliviero A, Delia Porta G, Peretti GM, Maffulli N. MicroRNA in osteoarthritis: physiopathology, diagnosis and therapeutic challenge. Br Med Bull. 2019;130(1):137–47.
20. Gargano G, Oliviero A, Oliva F, Maffulli N. Small interfering RNAs in human tendons. Br Med Bull. 2022;142(1):34–43.

Publisher’s Note
Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.