

Mesenchymal Stem Cell Therapy in Osteoarthritis and Rheumatoid Arthritis: A Systematic Review of Exosomal microRNAs

Seyyed Sina Hejazian^{1,2}, Seyyede Mina Hejazian³, Seyede Saba Mostafavi Montazeri⁴, Sima Abediazar³, Sepideh Zununi Vahed³, Abolfazl Barzegari⁵⁻⁷

¹Neurosciences Research Center, Tabriz University of Medical Sciences, Tabriz, Iran; ²Immunology Research Center, Tabriz University of Medical Sciences, Tabriz, Iran; ³Kidney Research Center, Tabriz University of Medical Sciences, Tabriz, Iran; ⁴Student Research Committee, Alborz University of Medical Sciences, Karaj, Iran; ⁵Department of Medical Biotechnology, Faculty of Advanced Medical Sciences, Tabriz University of Medical Sciences, Tabriz, Iran; ⁶Research Center for Pharmaceutical Nanotechnology (RCPN), Tabriz University of Medical Sciences, Tabriz, Iran; ⁷Laboratory for Vascular Translational Science, Nanotechnologies for Vascular Medicine and Imaging Team, Université Sorbonne Paris Nord, Villetaneuse, 93430, France

Correspondence: Sepideh Zununi Vahed, Kidney Research Center, Tabriz University of Medical Sciences, Tabriz, Iran, Email sepide.zununi@gmail.com; Abolfazl Barzegari, Department of Medical Biotechnology, Faculty of Advanced Medical Sciences, Tabriz University of Medical Sciences, Tabriz, Iran, Email abolfazl.barzegari@gmail.com

Summary: Osteoarthritis (OA) and rheumatoid arthritis (RA) are chronic diseases characterized by persistent inflammation and autoimmune responses that affect the joints and other organs. Scientific evidence indicates that the therapeutic effects of mesenchymal stem cells (MSCs) are mediated through the release of soluble factors and extracellular vesicles (EVs), particularly exosomes. The release of microRNAs from MSCs holds substantial potential for cell-free treatment in OA and RA.

Methods: A comprehensive search was conducted on Web of Science, PubMed, Scopus, and Google Scholar to identify relevant publications until 24 March 2024. The systematic review aimed at illuminating the current understanding of MSC-derived exosomal microRNAs, the origin of MSCs, potential mechanisms of action, and their therapeutic implications in managing OA and RA.

Results: A total of fifty-five articles (OA, n= 41 and RA, n= 14) were deemed eligible for inclusion in this study. Regardless of MSC origin, exosomal miRNAs could induce anti-inflammatory, protective, and chondroregenerative potential in in vitro and in vivo models of OA by targeting different signaling pathways. Fourteen studies have highlighted the role of MSC-derived exosomal miRNAs in modulating immune responses, reducing pro-inflammatory cytokine production, and potentially ameliorating synovial inflammation and joint symptoms associated with RA. To suppress joint inflammation and preserve cartilage, miR-140, miR-92a-3p, and miR-136-5p emerged as leading candidates for OA because they help restore the anabolic/catabolic balance and modulate key signaling pathways. For RA, the most effective candidates were miR-146a, miR-150-5p and miR-205-5p that target innate and adaptive immune signaling and synovioocyte activation, with NF-κB modulators such as miR-361-5p offering overlap across both diseases.

Conclusion: The mounting body of preclinical evidence supports that MSC-exosomal-miRNAs present a promising solution for OA and RA as a novel therapeutic strategy. However, human studies and more clinical trials are warranted.

Keywords: arthritis, extracellular vesicles, miRNAs, stem cells, MSCs, osteoarticular disease

Introduction

Osteoarthritis (OA) and rheumatoid arthritis (RA), the most common types of arthritis, are chronic inflammatory, autoimmune, and age-related osteoarticular diseases. They are characterized by the degeneration of cartilage and the inflammation of the synovial membrane, respectively. Both OA and RA involve inflammation and joint degeneration, although through different mechanisms.

Osteoarthritis is the most prevalent type of arthritis, involving over 250 million individuals worldwide.^{1,2} Currently, the prevalence of knee OA among the population aged over 40 is as high as 23%.³ At present, there are limited treatment options for knee OA, none of which can reverse cartilage degradation. Autologous chondrocyte implantation (ACI) is a

technique used for larger cartilage defects.⁴ In patients with more severe knee OA, total knee arthroplasty (TKA) is a good choice with remarkable outcomes; however, the artificial joints require revision 15 to 25 years after surgery.⁵

Rheumatoid arthritis is considered a systemic autoimmune disorder due to leukocyte influx into joints that produce inflammatory mediators and destroy bone and cartilage.⁶ The incidence of RA worldwide is 40 per 100,000 individuals per year, the prevalence is ~0.24%⁷ and men are significantly more exposed.⁸ Over the last decade, therapeutic options for RA have been improved; however, their serious complications and low efficacy in some patients remain unsolved. Some RA patients remain untreated because of their resistance or intolerant reactions.⁹ Thus, there is a necessity to develop other therapeutic strategies for RA.

Mesenchymal stem cell (MSC) therapy is a much newer therapy for patients with OA/RA, which practically fills the gap between conventional pharmaceutical and surgical therapies. This treatment is available in an intra-articular manner with a relatively low cost, but the risk of non-optimal transformation of the transplanted cells and hyper-inflammatory reactions is always there.¹⁰ This is why scientists have recently proposed the use of cell-free therapeutic strategies such as MSC-derived extracellular vesicles (EVs), especially exosomes (MSC-Exos), to enhance the ideal therapeutic impact against cartilage and bone diseases via selectively administering the specified factors.¹¹

It is worth mentioning that a part of the therapeutic effects of MSCs is attributable to EVs /exosomes, which carry selected cargo.¹² Given recent advancements, these exosomes are currently exploited as effective drug delivery strategies that can be modified to carry particular agents like small non-coding microRNAs (miRNAs; miR) as regulators of gene expression and various biochemical pathways.

Although research on EVs and MSC-Exos in joint diseases is increasing rapidly, existing reviews either address EV cargoes in general or focus on a single condition, such as OA or RA. However, miRNAs packaged within MSC-Exos represent a distinct and biologically active subset of EV cargo with demonstrated roles in cartilage repair, immunomodulation, and bone remodeling in preclinical models of both OA and RA.^{13,14} This specific area has not yet been systematically reviewed, creating a clear gap in the current literature. Numerous experimental studies have shown that transferring defined miRNA cargoes via MSC-derived exosomes can alter disease progression, highlighting their potential as diagnostic biomarkers and therapeutic candidates.¹⁵ However, published studies vary substantially in EV isolation, characterization, and miRNA analytic methods. Synthesizing evidence across OA and RA, therefore, offers a unique opportunity to identify miRNAs that function as shared mediators of joint pathology, as well as those specific to each disease. Such a cross-disease approach can reveal converging molecular mechanisms underlying cartilage degeneration and inflammation, which are not captured in single-disease reviews. In this review, the recent concepts of MSC derived exosomal-miRNAs are presented that can be applied to the treatment of joint disorders such as OA and RA.

microRNAs Carried by MSC-Derived Exosomes

The discovery of mRNAs and miRNAs being packed inside exosomes and shuttled between cells is a breakthrough, making these cargo-carrying components promising tools in various conditions. Since miRNAs participate in the regulation of inflammatory factors and chondrocyte homeostasis,¹⁶ they are important controllers in preserving a healthy joint, whereas Dicer knock-out mice are proof of this claim.¹⁷ Dicer is a crucial enzyme in the biosynthesis of miRNAs.

In both animal models and humans, dysregulated levels of miRNAs have been connected with osteoarticular diseases. Murata et al found that miRNA profiles of synovial fluid and plasma successfully discriminate RA from OA and can be used as diagnostic biomarkers. In synovial fluid of patients with RA, concentrations of miR-16, miR-146a, miR-223, and miR-155 were higher than those with OA significantly.¹⁸

Certainly, exosomal miRNAs have drawn greater attention than other molecules, given their functions in the regulation of gene expression. Besides, it is verified that a greater proportion of miRNAs is present in exosomes compared to their parent cells.¹⁹ More miRNA-profiling investigations have shown that the packaging of miRNAs into exosomes is done selectively. Accordingly, Guduric-Fuchs et al reported that subgroups of miRNAs, including miR-451, miR-142-3p, and miR-150, are preferentially wrapped inside exosomes.²⁰ Thus, miRNAs are thought to be the key functional components in target cells.

Exosomal miRNAs are preserved against endogenous RNase-based degradation by the exosomal membrane. Their capacity for survival in body fluids and cellular culture medium makes them suitable for intra-articular administration.

Further investigations have demonstrated that exosomal miRNAs are related to cartilage tissue development and homeostasis. A comparative microarray analysis between exosomes produced by undifferentiated human bone marrow-derived MSCs (hBM-MSCs) and cartilage-differentiating hBM-MSCs revealed that 35 miRNAs, consisting of miR-1290, miR-1246, miR-320c, miR-193a-5p, and miR-92a are upregulated in exosomes secreted from cartilage-differentiating hBM-MSCs.²¹ The same study also demonstrated that hBM-MSCs overexpressing exosomal miR-320c enhance the chondrocyte level of SOX9 (SRY-box transcription factor 9) and decrease MMP-13, leading to the expanded chondrogenesis of hBM-MSCs.²¹

Regarding the well-known cartilage regeneration role of miRNAs, miRNA (ie miR-155, miR-140, miR-136, miR-92a, miR-26a, and miR-9) -overexpressing exosomes indicate greater therapeutic perspective in the recipient cells in comparison to the naïve exosomes.^{13,22–25}

Methods

Search Strategy and Inclusion and Exclusion Criteria

The systematic review was carried out to examine the effects of exosomal miRNAs derived from MSCs on arthritis models, following the PRISMA guidelines. Various databases including Web of Sciences, PubMed, Scopus, and Google Scholar were searched with no time restriction, up to 24 March 2024 by a qualified librarian. The search strategy utilized terms and MeSH-based keywords, are presented in the [Supplementary Table S1](#). In this review, original research articles in English focusing on preclinical in vitro and animal models of arthritis, osteoarthritis, and rheumatoid arthritis utilizing MSCs secretome (exosomal miRNAs) were considered. Abstracts, review articles, non-English articles, retracted articles, and those with low quality were excluded. Two researchers independently reviewed records from the primary search (SMH and SSMM) with a third researcher resolving any discrepancies. Data extraction was performed by two researchers. The extracted information can be found in [Tables 1–4](#).

Assessment of the Quality of Studies

The quality of the studies included in the review was assessed based on the ARRIVE guidelines 2.0 and the risk of bias was evaluated using SYRCLE's risk of bias tool.

Results

Included Studies

The initial search yielded 447 articles, with 191 duplicates removed. One hundred and seven reports were screened through abstracts. Stem cell-based therapies, review articles, and irrelevant and retracted studies were excluded. Fifty-five articles were included in the review based on our criteria.^{13,15,21–49,51–73} The PRISMA flow diagram in [Figure 1](#) outlines the literature search process.

MSC Sources and Pre-Conditioning Methods to Obtain Engineered Exosomes in OA

Included OA articles were twelve in vitro,^{21,26–29,31–34,36–38} four in vivo,^{22,25,39,60} and twenty-five mixed in vivo+ in vitro studies, [Table 1](#). Different types of chondrocyte cell lines, including C5.18 and CHON-001 or isolated chondrocytes from cartilage of OA patients, were used in cell culture studies. IL-1 β or sodium iodoacetate was used to induce OA in most of these studies. The anterior cruciate ligament transection (ACLT) and destabilization of the medial meniscus (DMM) surgeries were commonly used surgical models to induce OA in animal studies. Additionally, other methods such as monosodium iodoacetate (MIA) injection⁴⁵ and collagenase VII^{25,56} administration were employed to create OA animal models.

Among OA studies, BM-MSCs were the most utilized cells.^{13,21,22,25,26,30,32,33,35,38–40,42,43,48,50,54,55,57,58} Beyond BM-MSC, S-MSCs,^{23,24,27,28,37,53,59,60} AD-MSCs,^{29,34,47,51,52} urine-derived MSCs,^{31,46} umbilical-cord MSCs (UC-MSCs),^{36,45} infrapatellar fat pad derived MSCs (IPFP-MSCs),⁴⁹ dental pulp-derived MSCs,⁴⁴ and undefined MSCs^{41,56} were utilized in OA studies. In most of the OA studies, MSCs were pre-conditioned with hypoxia,^{30,31} TGF- β ,^{41,48} curcumin,⁵⁵ fucoidan,⁵⁴ bFGF,⁵² quercetin,⁴³ or transfected with mimic/inhibitory miRNAs before EV/exosome isolation, [Table 1](#).

Table 1 In vitro/in vivo Osteoarthritis Models Applied MSC-Derived Exosomal microRNAs

No.	Authors	Year	Model			MSCs		MSCs Pre-Conditioning	Exosomes				Ref.
			Disease Model	Cell Line/Animals	Induction Model	Origin/Source of MSCs	Culture Media		Exosome Isolation	Characterization	Markers	Size	
1	Hu et al	2023	In vitro	Chondrocytes	IL-1 β	Rat BM-MSCs	DMEM Passage 3-5	Transfected with miR-140-3p mimics and inhibitors	Ultra-high-speed centrifugation	TEM, WB	CD9 ⁺ CD63 ⁺ TSG101 ⁺	-	[26]
2	Qiu et al	2021	In vitro	Chondrocytes	IL-1 β 10 ng/mL	hS-MSCs	RPMI1640	Transfected with miR-129-5p mimics and inhibitors	Ultracentrifuged at 100,000 \times g for 1h	TEM, WB	CD9 ⁺ CD63 ⁺ HSP70 ⁺	50-100 nm	[27]
3	Qiu et al	2024	In vitro	Cartilage cells	IL-1 β (10 ng/mL)	hS-MSCs	DMEM/F-12	Transfected with miR-485-3p mimics/inhibitors	Ultracentrifuged at 100,000 \times g for 1h	TEM, NTA, WB	CD81 ⁺ TSG101 ⁺	100 nm	[28]
4	Ragni et al	2020	In vitro	hADSCs	Fluid inflammatory conditions	hAD-MSCs	FBS-free DMEM Passage 5	Treated under inflammatory conditions, mimicking OA synovial fluid, the secretome of MSCs was used	Secretome (5mL) were diluted 1:2 with PBS and centrifuged at 100,000 \times g for 9 h	TEM, NTA, WB, flow cytometry	CD63 ⁺ CD81 ⁺ Low CD9	100 nm	[29]
5	Rong et al	2021	In vitro	Rat chondrocytes	IL-1 β	BM-MSCs rats	HG-DMEM Passage 3-5	Hypoxia, infected with a lentiviral miR-216a mimic/inhibitor vector	Ultra-15 centrifugal filter and centrifuged at 4000 \times g at 4 $^{\circ}$ C	TEM, NTA, WB	CD9 ⁺ TSG101 ⁺ CD63 ⁺ CD81 ⁺	50-150 nm	[30]
			In vivo	Female SD rats, aged 7–8 weeks, 200–250 g	DMM								
6	Wan et al	2022	In vitro	Chondrocyte CHON-001	-	Human urine-derived MSCs	DMEM Passage 3	Hypoxia, transfected with miR-26a-5p mimics/inhibitor	Centrifuged at 100,000 \times g for 2 h	TEM, WB, NanoSight tracking detector	CD63 ⁺ TSG101 ⁺ Syntenin ⁺ Calnexin ⁻	30-100 nm	[31]
7	Wang et al	2021	In vitro	Chondrocyte	IL-1 β	hBM-MSCs	α -MEM	-	10,000 g for 30 min, centrifuged at 100,000 \times g for 70 min	TEM, NTA, WB	CD9 ⁺ CD63 ⁺ HSP70 ⁺	100 nm	[32]
8	He et al	2020	In vitro	Mouse chondrocyte	LPS	Mouse BM-MSCs	DMEM	Transfected with or without miR-210	Centrifuged at 110,000 \times g at 4 $^{\circ}$ C for 1 h	TEM, NTA, WB	Alix ⁺ TSG101 ⁺ HSPA8 ⁺	30-150 nm	[33]
9	Li, Lin, and Liu	2023	In vitro	Chondrocyte	IL-1 β	Mouse AD-MSCs	DMEM	miR-93-5p mimic and miR-93-5p inhibitor	ExoQuick TC	TEM, NTA, WB	CD9 ⁺ CD81 ⁺ TSG101 ⁺	30-150 nm	[34]

10	Xu et al	2021	In vitro	Chondrocytes	IL-1 β (10 ng/mL)	BM-MSCs of femur rats	DMEM	Transfected with miR-326 mimics or inhibitors	ExoQuick-TC	TEM, NTA, WB	CD9 ⁺ , CD63 ⁺ , CD81 ⁺	~123 nm	[35]
			In vivo	Male SD rats (weighing 200–220 g)	Sodium iodoacetate (8%)								
11	Li et al	2021	In vitro	Articular cartilages	Obtained from 12 patients with knee arthroplasties	hUC-MSCs	DMEM Passage 3	Transfected by miR-100-5p mimic or inhibitor	Differential ultracentrifugation	Immunoblotting	CD9 ⁺ CD63 ⁺ CD81 ⁺	-	[36]
12	Zheng et al	2022	In vitro	Chondrocytes	IL-1 β (10 ng/mL)	hS-MSCs	DMEM Passage 3	Transfected by miR-212-5p mimic	CM was filtered (0.22 μ M filters), centrifuged at 4000 \times g into ~200 μ L and further ultracentrifuged at 100,000 \times g 1 h, 4 $^{\circ}$ C	TEM, DLS	CD105 ⁺ CD44 ⁺ SCA-1 ⁺	30-150 nm	[37]
13	Dong, Li et al	2021	In vitro	Chondrocytes	IL-1 β (10 ng/mL)	BM-MSCs from femur of SD rats	DMEM Passage 1-3	Transfected with miR-127-3p (NC) inhibitor	Supernatant was centrifuged at 4 $^{\circ}$ C at 2000 \times g for 15 min and then was centrifuged at 4 $^{\circ}$ C at 100,000 \times g for 1 h. The precipitate was resuspended in PBS and centrifuged at 100,000 \times g for 1 h at 4 $^{\circ}$ C	TEM, WB, NTA	CD29 ⁺ CD44 ⁺ CD34 ⁺ CD45 ⁺	30-120 nm	[38]
14	Mao et al	2018	In vivo	6-week-old female C57B/L10 mice	12 U collagenase VII in 8 μ L saline, Clostridium histolyticum	hBM-MSCs	α -MEM Passage 5	Transfected with miR-92a-3p mimic or inhibitor (40 nM)	3000 \times g (30 min) and 10,000 \times g (30 min). Then 64,000 \times g for 110 min using an SW28 rotor. Resuspended in 0.32 M sucrose and centrifuged at 100,000 \times g for 1 h	TEM, WB	CD9 ⁺ CD63 ⁺ CD81 ⁺ HSP70 ⁺	50-150 nm	[25]
15	Sun et al	2019	In vitro	Human OA articular cartilage samples	Obtained from three male patients	hBM-MSCs	α -MEM Passage 3	Transfected with miR-320c mimic or a nonspecific control (50 nM)	Filtered (0.22 μ m), centrifuged at 3000 \times g for 15 min at 4 $^{\circ}$ C. 5 mL supernatant incubated overnight with ExoQuick reagent (1 mL) and centrifuged at 1500 \times g for 30 min at 4 $^{\circ}$ C	NTA, TEM, WB	CD9 ⁺ CD63 ⁺	50-200 nm	[21]
16	Wang & Xu	2021	In vivo	12-week-old SD rats	ACLt and DMM	BM-MSCs 3-5-day-old newborn rats	α -MEM	Inhibitor negative control (INC), miR-135b inhibitor (miR-135b), miR-135b mimic, and mimic negative control	Exosome extraction kit (Genesee, Guangzhou, China)	TEM, NTA, WB	CD29 ⁺ CD44 ⁺ CD90 ⁺ low CD11b low CD34 low CD45	100 nm	[39]

(Continued)

Table I (Continued).

No.	Authors	Year	Model			MSCs		MSCs Pre-Conditioning	Exosomes				Ref.
			Disease Model	Cell Line/Animals	Induction Model	Origin/Source of MSCs	Culture Media		Exosome Isolation	Characterization	Markers	Size	
17	Huang, et al	2021	In vivo	Adult 9-week-old C57BL/6 mice	Mouse right knee joint was exposed by the medial para-patellar approach. The patella was dislocated, and the knee was completely flexed. The anterior cruciate ligament was then cut off by a micro-scissor.	BM-MSCs from 5 C57BL/6 mice aging 4 week	DMEM/F12 Passage 4	miR-206 mimic, miR-206 inhibitor and their NCs	ExoQuick-TC exosome precipitation liquid (ExoQuick-TC Kit; System Biosciences)	TEM, NTA, WB	CD29 ⁺ CD90 ⁺ CD63 ⁺ Alix ⁺	95.4 ± 18.9 nm	[40]
			In vitro	Knee joint samples	5 remaining mice in the OA group								
18	Wang, Xu, and Xu	2018	In vitro	Rat chondrocyte line (C5.18 cell)	-	MSCs from male SD rats	DMEM	TGF-β1 (10 ng/mL) for 24 h	Exosome extraction kit (Geneseeed, Guangzhou, China)	TEM, WB	CD81 ⁺ ALIX ⁺		[41]
			In vivo	24 male 12-week-old SD rats	DMM and ACLT								
19	Wang et al	2021	In vitro	Primary chondrocytes	Extracted from fragmented cartilage tissue of OA patients	hS-MSCs	DMEM Passage 1	Transfected with, miR-155-5p mimic, and miR-155-5p inhibitor	-	NTA, TEM, WB	CD81 ⁺ CD63 ⁺ TFIIB ⁺	100-120 nm	[23]
			In vivo	20 specific SPF BALB/C mice	4 °C cold water stimulation for 2 h twice a day								
20	Jin et al	2020	In vivo	6-week-old 80 male SPF SD rats	ACLT/MCL transection	BM-MSCs from bilateral femora and tibia	L-DMEM PASSAGE 2	Transfected with miR-9-5p mimics, inhibitors, or their NCs (100 pmol)	Ultracentrifugation	WB, TEM, flowcytometry	CD9 ⁺ TSG101 ⁺ CD63 ⁺	40-100 nm	[22]
21	Zhang, Qi, et al	2023	In vitro	Murine chondrocyte of newborn C57BL/6 mice	IL-1β (10 ng/mL)	BM-MSC from femur and tibia of C57BL/6 mice (4 weeks old)	DMEM Passage 3	-	CM was centrifuged at 3000 ×g for 15 min, followed by another 30 min of centrifugation at 10,000 ×g. Then the supernatant was re-centrifuged at 4000 ×g for 30 min at 4 °C. ExoQuick reagent was added at a 1:5 volume and incubated at 4 °C for 12 h. Exosomes were pelleted by centrifuging at 1500 ×g for 30 min at 4 °C.	NTA, TEM, WB	HSP70 ⁺ CD63 ⁺ TSG101 ⁺	30-150 nm	[42]
			In vivo	C57BL/6 mice	DMM surgery								

22	Dong et al	2024	In vitro	Human chondrocytes CHON-001	IL-1 β for 24 h	BM-MSCs were provided with COBIER	RPMI-1640 Passage 1	-transfected with NC or miR-124-3p agomir - incubated with 20 mM Quercetin for 24 h	BM-MSCs were centrifuged for 15 min at 300 \times g, 2000 \times g for 15 min, 10,000 \times g for 30 min, and the supernatants were ultracentrifuged for 70 min at 120,000 \times g	NTA, TEM, WB	TSG101 ⁺ CD9 ⁺ CD81 ⁺	30-150 nm	[43]
			In vivo	12-week-old 12 male Wistar rats	DMM surgery of the right knee								
23	Lin et al	2021	In vitro	Cartilage cells	IL-1 β (10 ng/mL)	Dental pulp stem cells (dPSCs)	α -MEM Passage 3	Overexpressed miR-140-5p	Culture supernatants were centrifuged at 3000 \times g for 30 min and then at 10,000 \times g for 30 min at room temperature. Pellets centrifugation was at 64,000 \times g for 110 min at 4°C and washed once with 0.32 M sucrose	NTA, TEM, WB	CD9 ⁺ CD63 ⁺	134 \pm 29 nm	[44]
			In vivo	~8-week-old SPF male SD rats	50 μ L iodoacetic acid (25 mg/mL) was administered to the double knee joints								
24	Liu et al	2023	In vitro	Chondrocyte cells form articular cartilage of male 5-day-old SD rats	IL-1 β (10 ng/mL) for 24 h	hUC-MSCs	α -MEM Passage 1	Mix 0.5 μ mol miR-223 mimics with 10 μ g of total EVs	Ultracentrifugation	NTA, TEM, WB	CD81 ⁺ Tsg101 ⁺ Alix ⁺ Calnexin ⁻	Average of 142.9 nm	[45]
			In vivo	8-week-old SD rats	Through the infrapatellar ligament, 50 μ L MIA at 20 mg/mL was injected intraarticularly								
25	Liu et al	2022	In vitro	Primary chondrocytes isolated from part of the cartilage	IL-1 β (10 ng/mL) for 24 h	hU-MSCs	Passage 3	Transfected with miR-140-5p	Ultracentrifuged	TEM, NTA, WB	CD63 ⁺ CD9 ⁺ CD81 ⁺ calnexin ⁻	50-150 nm	[46]
			In vivo	12-week-old female SD rats	ACLT and DMM of the right knee								
26	Meng et al	2023	In vitro	Primary articular chondrocytes	Separated from knee articular cartilage fragments harvested from the femoral condyles and tibial plateaus of 8-week-old SD female rats	Rat AD-MSCs	DMEM/F12 Passage 3-5	Tropoelastin (TE) for 48 h	1 mL of CM was mixed with 500 μ L total exosome isolation reagent, incubated overnight at 4 °C, and the mixture was centrifuged at 10,000 \times g for 1 h at 4 °C. The exosome pellet was resuspended in PBS	TEM, NTA, WB	TSG101 ⁺ CD81 ⁺	50-200 nm	[47]
			In vivo	Male ~12-week-old SPF SD rats	Transecting the ACLT								

(Continued)

Table I (Continued).

No.	Authors	Year	Model			MSCs		MSCs Pre-Conditioning	Exosomes				Ref.
			Disease Model	Cell Line/Animals	Induction Model	Origin/Source of MSCs	Culture Media		Exosome Isolation	Characterization	Markers	Size	
27	Sun et al	2022	In vitro	hBM-MSCs	TGFβ3 (50 ng/mL) every 2–3 d for 7 d	hBM-MSCs isolated from biopsies of healthy clinical donors	DMEM Passage 3	-TGFβ3 (50 ng/mL) for 7 days before EV collection -transfected with miRNA-455 mimics, inhibitors, or negative control	Collected CM (500 mL) was centrifuged at 300 ×g for 10 min. The resulting supernatant was filtered (0.4-μm cell strainer and 0.22-μm bottle top filter). The suspension was subjected to a tangential flow filtration system (TFF) with a 300 kDa MWCO capsule.	TEM, DLS, NTA, WB	CD9 ⁺ CD63 ⁺ CD81 ⁺ Alix ⁺	50-200 nm	[48]
			In vivo	Adult male rats	ACLT								
28	Wu et al	2019	In vitro	Chondrocytes	IL-1β (10ng/mL)	MSC of Infrapatellar fat pad-derived from OA patients	DMEM/F12 Passage 1	Unpretreated or pretreated with tropoelastin (TE)	-ExoQuick™ (EQ) reagent kit (SBI) and ultrafiltration	TEM, WB, Flowcytometry	CD81 ⁺ CD9 ⁺ CD63 ⁺	30-150 nm	[49]
			In vivo	9-week-old male C57BL/6 mice	Surgical DMM of the right knee								
29	Ye et al	2022	In vitro	Chondrocytes from the knee joint of 4 C57BL/6 mice	IL-1β (10 ng/mL) for 24 h	Mouse BM-MSCs	DMEM Passage 1	Mimic NC/miR-3960 mimic, inhibitor-NC/miR-3960 inhibitor	Ultracentrifuged	WB, NTA	CD9 ⁺ CD63 ⁺ TSG101 ⁺ Calnexin ⁻	30-150 nm	[50]
			In vivo	32 C57BL/6 J male 8-week-old mice	Transversely Sheared medial collateral ligament at the joint junction and the incised capsule at the upper and lower edges of the meniscus								
30	Meng et al	2023	In vitro	Chondrocytes from ADSCs from male 4-week-old SD rats	IL-1β	AD-MSCs from male SD rats 4 weeks old	(DMEM)-F12	miR-429 mimics, inhibitor, miR-NC (negative control), and pcDNA3.1- FEZ2	CM was centrifuged at 4 °C for 300 ×g for 10 min, followed by 20-min centrifugation at 4 °C and 2000 ×g, and 45-min centrifugation at 4 °C and 11,000 ×g. The supernatant was centrifuged at 110,000 ×g at 4 °C for 90 min, the precipitate resuspended in 1 mL 1× PBS, and centrifuged at 110,000 ×g at 4 °C for 70 min	TEM, WB, Granulometric analysis	CD81 ⁺ CD63 ⁺	30-150 nm	[51]
			In vivo	60 SPF SD male 6-week-old rats	Intravenously injected with 0.1 mL 8% sodium iodoacetate solution once daily for 7 consecutive days								

31	Zhao et al	2023	In vitro	Primary articular chondrocytes from newborn SD rats	IL-1 β for 24 h	Subcutaneous fat (SC) MSCs from adipose tissue	DMEM/F12 Passage 2	bFGF (10 ng/mL) for 24 h	Ultracentrifuged	NTA, TEM, WB	CD9 ⁺ CD81 ⁺ Alix ⁺ TSG101 ⁺ Calnexin ⁻	70-150 nm	[52]
			In vivo	6-week-old male SD rats	DMM and ACLT surgery								
32	Wang et al	2020	In vitro	36 articular cartilage tissues	Obtained from OA subjects	hS-MSCs isolated from the synovium of human knee joints	Passage 3	miR-31 mimic, sh-PTTG1-1, and shPTTG1-2	ultraCentrifuged	DLS, TEM, NTA, WB	CD81 ⁺ CD9 ⁺ CD63 ⁺ Calnexin ⁻	95.01 \pm 35.91 nm	[53]
			In vivo	20 male 8-week-old C57 mice	DMM and ACLT								
33	Lou et al	2023	In vitro	Rat chondrocyte	IL-1 β (10 ng/mL)	BM-MSCs isolated from the femurs of 4-week-old male SD rats	DMEM Passage 1-3	Fucoidan for 48 h	Ultracentrifugation	DLS, TEM, WB	CD9 ⁺ CD63 ⁺ CD81 ⁺ TSG101 ⁺ Calnexin ⁻	144.0–156.7 nm	[54]
			In vivo	48 male 10-week-old SD rats	ACLT and DMM								
34	Qiu, Xu, Yi, and Ha	2020	In vitro	Primary chondrocytes	IL-1 β (10 ng/mL)	BM-MSCs	DMEM medium	Curcumin	Supernatant was centrifuged at 300 \times g for 10 minutes followed by 15 minutes of centrifugation at 2000 \times g. Then, the supernatant was filtered (0.22 μ m) followed by 70 minutes of ultra-filtration at 100000 \times g, which were then re-suspended in PBS	MTT, WB, TEM	CD9 ⁺ CD63 ⁺ CD81 ⁺	50-150 nm	[55]
			In vivo	Mice models	OA operation								
35	Liu et al	2018	In vitro	Chondrocytes	Isolated from the knee joint of C57BL/6 mice	hMSCs	DMEM/F12 Passage 3	Lentiviral vectors that stably overexpressing KLF3-AS1, lentiviral vectors that stably silencing KLF3-AS1, siRNA targeting GIT1 (si-GIT1), miR-206 mimic, and their corresponding controls	Centrifuged at 12,000 \times g for 1 h.	TEM, flowcytometry	TSG101 ⁺ CD63 ⁺		[56]
			In vivo	Adult SPF 9-week-old C57BL/6 mice	Type II collagenase solution was dissolved in sterile PBS (pH 7.4), filtered and then injected into the knee joint cavity								

(Continued)

Table I (Continued).

No.	Authors	Year	Model			MSCs		MSCs Pre-Conditioning	Exosomes				Ref.
			Disease Model	Cell Line/Animals	Induction Model	Origin/Source of MSCs	Culture Media		Exosome Isolation	Characterization	Markers	Size	
36	Xia, Wang, Lin, and Wang	2021	In vitro	Chondrocytes from cartilage excised from the subchondral bone		BM-MSCs from the femur of trauma OA patients	DMEM/F-12	Mimics, inhibitors, and the negative controls of miR-125a-5p	Ultrafiltrate	WB, TEM	CD63 ⁺ CD9 ⁺	50–150 nm	[57]
			In vivo	48 healthy male 5–8-week-old C57BL/6 mice	Applying a single mechanical load (1 mM/s to 12 N) to the ankle joint causes the tibia to move forward relative to the femur and extend the anterior cruciate ligament beyond the point of failure.								
37	Tao et al	2021	In vitro	Chondrocytes from articular cartilage tissues	IL-1 β 10 (ng/mL) for 24 h	hBM-MSCs	DMEM Passage 1	Load the exosomes with miR-361-5p and miR-NC via electroporation	ExoQuick-TC reagent	-	-	-	[58]
			In vivo	60 Wistar rats	A parapatellar incision was made to expose the patella of the knee, and the patella was dislocated laterally. The anterior cross-ligament was exposed by bending the knee as far as possible, and the anterior cruciate ligament was cut off under direct vision								
38	Chen et al	2020	In vitro	Cartilage dissected from the subchondral bone	20 cases of cartilage tissues from patients with traumatic OA of the knee and 18 cases of knee cartilage tissues from patients with post-traumatic amputation	Femur BM isolated from the traumatized patients	MesenGro [®] hMSC medium	Lentiviral vectors containing ELF3 overexpression	Ultracentrifuged	NTA, TEM, WB	CD63 ⁺ CD9 ⁺ CD81 ⁺ Alix ⁺ TSG101 ⁻	50-150 nm	[13]
			In vivo	48 healthy male 5–8 week-old C57BL/6 mice	A single mechanical load was applied to the ankle joint ((1 mM/s to 12 N), causing the tibia to move forward relative to the femur and extended the anterior cruciate ligament beyond the point of failure.								

39	Lu et al	2021	In vitro	Human chondrosarcoma SWI353 cells	IL-1 β (10 ng/mL) for 24 h	Synovial tissues (20–50 mg wet weight) from aseptic synovial biopsy	DMEM Passage 3	LV interference vectors of miR-26a-5p	Ultracentrifuged	TEM, WB, NTA	CD63 ⁺ CD9 ⁺ Calnexin ⁻	100-120 nm	[59]
			In vivo	36 adult male SD rats	DMM and ACLT								
40	Tao et al	2017	In vitro	Chondrocytes from human cartilage	Obtained from the resected lateral femoral condyle of patients, aged from 45 to 55 years old	Synovial membranes aseptically harvested from the knee joint of trauma 5 patients, aged from 30 to 35 years old	Passage 5	Overexpression of miR-140	Ultracentrifuged	DLS, TEM, WB	CD63 ⁺ CD9 ⁺ CD81 ⁺ Alix ⁺	30-150 nm	[24]
			In vivo	Male ~12 weeks old SD rats	DMM and ACLT								
41	Kong et al	2023	In vivo	8 weeks old SD rats	DMM	Synovial membrane	Passages 3–4	Mimic and miR-320c mimic transfection	HiefTM Quick Exosome Isolation Kit	-	-	-	[60]

Abbreviations: ACLT, anterior cruciate ligament transection; ADSCs, adipose-derived stromal cells; BMSCs, Bone mesenchymal stem cells; CIA, collagen-induced arthritis; DMM, destabilization of the medial meniscus; hSMSCs, human synovial microvascular endothelial cells; hUSCs, human urine-derived stem cells; hUC-MSCs, Human umbilical cord-derived mesenchymal stem cells; MCL, medial collateral ligament; NTA, nanoparticle tracking analysis; OA, Osteoarthritis; RA, Rheumatoid arthritis; RA-FLSs, RA-fibroblast-like synoviocytes; SD, sprague dawley; SPF, specific-pathogen free; TEM, transmission electron microscopy; WB, Western blotting.

Table 2 In Vitro/in Vivo Rheumatoid Arthritis Models Applied MSC-Derived Exosomal microRNAs

No.	Authors	Year	In vitro/in vivo Models			MSCs		MSCs Pre-Conditioning	Exosomes				Ref.
			Disease Model	Cell Line/ Animals	Induction Model	Origin/ Source of MSCs	Culture Media		Exosome Isolation	Characterization	Markers	Size	
1	Tavasolian et al	2020	In vitro	Splenocytes	Isolated from CIA mice	AD-MSCs derived from CIA mice	DMEM	Genetically modified to overexpress miR-146a/miR-155	Centrifuge at 300 ×g for 15 min and supernatant was filtered (0.22 μm). Centrifuge at 2600 ×g for 15 min, centrifuge at 10,000 ×g for 70 min and centrifuge at 110,000 ×g for 2 h	TEM, SEM, Bradford Assay, DLS	CD9 ⁺ CD63 ⁺	30-150 nm	[15]
2	Meng et al	2020	In vitro	FLSs MH7A	-	hBM-MSCs	DMEM	Transfected by miR-124a overexpression vector; 2–3 days	Invitrogen kit	TEM, DLS, WB, flow cytometry	CD9 ⁺ CD63 ⁺	100 nm	[61]
3	Su et al	2021	In vitro	FLSs MH7A	-	hBM-MSCs	DMEM Passages 3-5	Transfected by HAND2-AS1 over-expression vector	Ultracentrifugation	TEM, NTA, WB	TSG101 ⁺ CD63 ⁺ CD81 ⁺ GRP94 ⁻	120 nm	[62]
4	Chen et al	2018	In vitro	FLSs MH7A	-	Mice BM-MSCs	DMEM Passages 3–6	Transfected with LAN-modified miR-150-5p	ExoQuick-TC Kit	TEM, NTA, WB	CD9 ⁺ CD63 ⁺	100 nm	[63]
			In vivo	7–9-week-old DBA/1 Male mice	CIA								
5	Huang et al	2022	In vitro	RAW264.7	LPS (1 μg/mL) for 12 h + ATP (5 mM) for 30 min	BM-MSCs from the bone marrow of femur and tibia	DMEM After Passage 3	-	Centrifugation at 3000 ×g for 15 min and then 0.5 vol of exosomes isolation reagent was added and incubated 4°C overnight. The mixture was centrifuged at 10,000 ×g for 1 h at 4°C and the pellets were resuspend in PBS	TEM, WB	CD81 ⁺ CD63 ⁺	~100 nm	[64]
			In vivo	40 male SPF SD rats	Subcutaneously injected with 0.1 mL of complete freund's adjuvant (CFA) to right posterior sole								
6	Lin et al	2023	In vitro	Human RA-FLS	LPS (1 μg/mL)	S-MSCs	DMEM Passage 4-7	Transfected by miR-433-3p mimics	Differential centrifugation	TEM, NTA, WB	CD9 ⁺ CD63 ⁺	-	[65]
			In vivo	8–10-week-old Male DBA/1J mice	100 μg bovine type II collagen (CII) via intradermal injection at the base of tail. On Day 21 after primary immunization, the mice received a booster shot of 100 μg CII								

7	Huang et al	2022	In vitro	Rat RA-SFs		hUC-MSCs	DMEM Passage 3–5	Transfected with miR-140-3p inhibitor or mimic	ExoQuick TC kit	TEM, NTA, WB, flow cytometry	CD81 ⁺ Alix ⁺	91 ± 33 nm	[66]
			In vivo	6-week-old Wistar Male rats	Seeded with bovine collagen II (100 µL) + complete Freund's adjuvant (100 µL) + mycobacterium tuberculosis								
8	Mi et al	2024	In vitro	RA SFs	-	hUC-MSCs	DMEM Passage 3–5	AGO2 knocking down in hUC-MSC (hUC-MSC KD-AGO ²)	Gradient ultra-high-speed centrifugation	TEM, NTA, WB	CD9 ⁺ CD63 ⁺ TSG101 ⁺	122 nm	[67]
			In vivo	6–8-week-old Wistar Female rats	CIA								
9	Meng and Qiu	2020	In vitro	hRA-FLSs	-	hBM-MSCs	DMEM Passage 3–7	Transfected with miR-320a mimic and mimic-NC plasmids	100,000 × g for 90 min	TEM, NTA, WB, flow cytometry	CD9 ⁺ CD63 ⁺ calnexin ⁻	30–150 nm	[68]
			In vivo	7-week-old C57BL/6 male mice	Collagen-induced arthritis model								
10	Ma et al	2022	In vitro	Mouse FLSs	IL-1β	Mice BM-MSCs	α-modified Eagle medium passages 3–6	Chondrogenic BM-MSCs	Total Exosome Isolation Reagent+ 10000 g for 70 min at 4°C	TEM, Laser particle size analyzer, WB	CD63 ⁺ CD9 ⁺	100 nm	[69]
			In vivo	8-week-old C57BL/6 Male mice	CIA								
11	Li et al	2021	In vitro	Mouse FLSs	-	Mouse BM-MSCs	DMEM Passage 3	miR-21 silenced in BMSCs	Ultracentrifugation	TEM, WB, DLS	CD63 ⁺ TSG101 ⁺ Calnexin ⁻	30–100 nm	[70]
			In vivo	96 BALB/c male mice	CIA								
12	Zhang et al	2023	In vitro	HSMEC	-	hBM-MSCs	DMEM Passage 3	Treated BMSCs with miR-378a-5p mimic or inhibitor	4°C and 100,000 g overnight	TEM, NTA, WB	CD63, and CD9	40–200 nm	[71]
			In vivo	8–12-week-old DBA/1 mice	CIA								
13	Wu et al	2021	In vitro	Rat RA-FLS	-	BM-MSCs	DMEM Passage 3	Transfected with the miR-34a inhibitor and miR-negative control	Centrifuged at 1. 300×g, 10 min, 2. 2000×g, 30 min 3. 10,000× g, 1 h	TEM, NTA, WB	CD81 ⁺ CD63 ⁺ calnexin ⁻	93 nm	[72]
			In vivo	6-week-old 30 Wistar male rats	Injected with Freund's complete adjuvant (0.1 mL/100 g) intradermally through the left hindlimb toes								
14	Wu et al	2022	In vivo	Mice model	-	BM-MSCs	-	-	-	-	-	-	[73]

Abbreviations: ADSCs, adipose-derived stromal cells; BMSCs, Bone marrow mesenchymal stem cells; CIA, Collagen-induced arthritis; DMEM, dulbecco's modified eagle medium; hS-MSCs, human synovial mesenchymal stem cells; hUC-MSCs, Human umbilical cord-derived mesenchymal stem cells; NTA, nanoparticle tracking analysis; RA, Rheumatoid arthritis; RA-FLSs, RA-fibroblast-like synoviocytes; SD, sprague dawley; SPF, specific-pathogen free; TEM, transmission electron microscopy; WB, Western blotting;

Table 3 Preclinical Studies Applied MSC-Derived Exosomal microRNAs for Osteoarthritis-Additional Information

No.	Author	Disease Model	Groups/Treatments	Concentration/Volume	Treatment/Delivery Route	Frequency	Follow Up	microRNAs	Target Gene	Functional Outcome	Ref.
1	Hu et al, 2023	In vitro	NC BM-MSC-Exo miRNA mimics control OE-miR-140- 3p inhibitor-miR-140-3p	100 µg/mL	Exosome incubated with chondrocytes	At h 0	-	miR-140-3p	HIF-1α	↑Chondrogenic differentiation ↓pathogenesis in OA	[26]
2	Qiu et al, 2021	In vitro	Control miR-NC-Exo miR-129-5p-Exo	-	Exosome incubated with IL-1β-induced chondrocytes	At h 0	24 h	miR-129-5p	HMGB1	↓ inflammatory response and ↓apoptosis of chondrocytes	[27]
3	Qiu et al, 2024	In vitro	Control OA Exos Exos + oe-NC Exos + oe-NRP-1	Exosomes in PBS (100 µL)	Treated with cartilage cells or mouse primary chondrocytes (10 ⁶ cells/well)	At h 0	2 h	miR-485-3p	NRP1-PI3K/Akt pathway	↓inflammation, ECM degradation and apoptosis of osteoarthritic cartilage cells ↓OA	[28]
4	Ragni et al, 2020	In vitro	Studying the MSCs-secreted factors and EV-embedded miRNAs under inflammatory conditions	-	MSCs treated with inflammatory conditions	At h 0	-	miR-24-3p	<i>PI6INK4A</i>	Regulates aging and cartilage catabolism	[29]
								miR-222-3p	<i>HDAC4</i>	↓cartilage degradation	
								miR-193b-3p	<i>TGFB2-3</i> <i>MMP19</i>	↓inflammation	
5	Rong et al, 2021	In vitro	PBS Normoxia-sEVs Hypoxia-sEVs	150 µg/mL	sEVs incubated with chondrocytes	At h 0	24 h	miR-216a-5p	JAK2/ STAT3	↑proliferation and migration and ↓apoptosis of chondrocytes	[30]
		In vivo	Normoxia-sEVs Hypoxia-sEVs miR-NC-Hypo-sEVs, silenced miR-216a-5p- Hypo-sEVs	200 µg of total protein of sEVs precipitated in 200 µL PBS	Injected into the joint cavity	At d 0	After 4 w			sEV miR-216a involved in the OA cartilage repair process	
6	Wan et al, 2022	In vitro	PBS Normoxia-sEVs Hypoxia-sEVs inhibitor-NC-Hypo-EVs miR-inhibitor-Hypo-EV	150 µg/mL	EVs incubated with chondrocytes	At hour 0	24 h	miR-26a-5p	PTEN	↑migration and proliferation of chondrocytes	[31]
7	Wang et al, 2021	In vitro	Control IL-1β IL-1β + Exo	0, 1, 5, 10, 20, and 50 µg/mL	Exosomes incubated with 10 ⁴ Chondrocytes/well	24, 48, and 72 h	72 h	miR-6515-5p	LYRM4- ASI GRPR- miR-6515- 5p	Anti-inflammatory effects	[32]

8	Lei He et al, 2020	In vitro	BM-MSCs-Exos BM-MSCs-vector-Exos BM-MSCs-210-Exos	-	-	-	-	miRNA-210	Tnfrsf21 NF-κB pathway	↑proliferation ↓apoptosis of chondrocytes	[33]
9	Li et al, 2023	In vitro	IL-1β + Exo IL-1β + mimic NC IL-1β + miR-93-5p inhibitor IL-1β + inhibitor NC IL-1β + miR-93-5p inhibitor	-	AD-MSC-exos co-incubated with the IL-1β-treated chondrocytes	-	24 h	miR-93-5p	ADAMTS9	↓apoptosis and autophagy through PI3K/AKT/mTOR signaling	[34]
10	Xu & Xu, 2021	In vitro	Normal chondrocyte OA chondrocyte Normal chondrocyte + BMSC-Exo	10 μg/mL	Exosomes incubated with chondrocytes	12 h	1, 3, 5, and 7 d	miR-326	HDAC3 STAT1 NF-κB p65	↓pyroptosis of cartilage and chondrocytes	[35]
		In vivo	Sham, n= 6 OA, n= 6 OA+ BMSC-Exo group, n= 6 OA+ BMSC-miR-326-Exo, n= 6	40 μg/100 μL	Intra-articular injection	Once a week	6 w				
11	Li et al, 2021	In vitro	Control 20% Cyclic strain + Vehicle 20% Cyclic strain + Exo 20% Cyclic strain + Exo + GW (20mM)	50 μg/mL	Exosomes incubated with chondrocytes	24 h	-	miR-100-5p	NOX4	↓ROS production and cell apoptosis in articular chondrocytes	[36]
12	Zheng et al, 202	In vitro	NC IL-1β IL-1β + Exo NC + Exo-miR-212-5p	2 μg	Exosomes incubated with chondrocytes for	48 h	-	miR-212-5p	ELF3	↓Chondrocyte Degeneration and inflammation	[37]
13	Dong et al, 2021	In vitro	1. Control chondrocytes 2. OA chondrocytes 3. OA chondrocytes + Exo 4. OA chondrocytes + Exo-NC 5. OA chondrocytes + Exo-inhibitor 6. OA chondrocytes + miR-127-3p NC mimic 7. OA chondrocytes + miR-127-3p mimic 8. OA chondrocytes + Exo + pcDNA empty plasmid 9. OA chondrocytes + Exo + pcDNA-CDH11 plasmid	20 μg/mL	Exosomes incubated with rat chondrocytes	24 h	6, 12, or 24 h	miR-127-3p	CDH11	↓Wnt/β-catenin pathway activation ↓chondrocyte damage	[38]

(Continued)

Table 3 (Continued).

No.	Author	Disease Model	Groups/Treatments	Concentration/Volume	Treatment/Delivery Route	Frequency	Follow Up	microRNAs	Target Gene	Functional Outcome	Ref.
14	Mao et al, 2018	In vivo	Sham, n= 10 OA, n = 10 MSC-Exos, n= 10 MSC-miR-92a-3p-Exo, n= 10	15 μ L exosome (500 μ g/mL)	Injection	Days 7, 14, and 21	28 d	miR-92a-3p	WNT5A	\uparrow production of cartilage and homeostasis	[25]
15	Sun et al, 2019	In vitro	Control hBMSC-Exo hBMSC-320c-Exo	10^8 /mL	Incubation	24 or 72 h	0, 24, 72, 120, and 168 h after the scratch	miR-320c	MMP13	\uparrow OA chondrocyte proliferation	[21]
16	Wang & Xu, 2021	In vivo	Sham group, n= 6 OA group, n= 6 OA + BMSC-Exo ^{PBS} , n= 6 OA + BMSC- Exo, TGF- β^1 n= 6 OA + INC-BMSC-Exo, TGF- β^1 n= 6 OA + miR-135bl-BMSC-Exo, TGF- β^1 n = 6	100 μ L exosome (10^{11} particles/mL)	Intraarticular cavity injection	-	-	miR-135b	MAPK6	\downarrow cartilage injury \uparrow M2 synovial macrophage polarization	[39]
17	Huang et al, 2021	In vitro	OA group Exo group mimic NC-Exo group miR-206 mimic-Exo group mimic NC group miR-206 mimic group sh-Elf3-NC group sh-Elf3 group miR-206 mimic + overexpressed (oe)-Elf3 group		Incubation		72 h	miR-206	Elf3	\uparrow osteoblast proliferation and differentiation	[40]
		in vivo	sham group, n = 10 OA group, n = 15 Exo group, n = 10 miR-206 mimic NC-Exo group, n = 10 miR-206 mimic-Exo group, n = 10 miR-206 inhibitor-Exo group, n = 10	100 μ L exosomes twice/w (10^{11} exosome particles/mL)	Injection of 100 μ L exosomes twice a week (10^{11} exosome particles/mL) into 10 knee joints from 5 mice	injected into 10 knee joints from 5 mice	8 w				

18	Wang et al, 2018	In vitro	TGF- β 1 + NC+si-control group TGF- β 1 + miR-135b inhibitor +si-control group TGF- β 1 + NC+si-Sp1 group TGF- β 1 + miR-135b inhibitor+si-Sp1 group	10 μ g/mL MSC-exosomes	Co-culture	72 h		miR-135b	Sp1	\uparrow chondrocyte proliferation	[41]
		In vivo	(1) OA+MSC-exosome group, n = 6 (2) OA+TGF- β 1 + MSC-exosome group, n = 6 (3) OA+TGF- β 1 + MSC-NC-exosome group, n = 6 (4) OA+TGF- β 1 + MSC-miR135b inhibitor-exosome group, n= 6	100 μ L exosome (10^{11} exosome particles/mL)	Articular cavity injection	-	12 w				
19	Wang et al, 2021	In vitro	Control SMSC-Exos SMSC-155-5p-Exos inhibitor SMSC-155-5p-Exos	10×10^{11} particles/mL	Incubation	36 h	96 h	miR-155-5p	Runx2	\uparrow proliferation and migration of chondrocytes \downarrow apoptosis	[23]
		In vivo	(1) Normal group, n = 10 (2) OA group n= 10 (3) OA + SMSC-Exo group, n= 10 (4) OA + SMSC-155-5p-Exo group, n= 10	30 μ L exosome (10^{11} particles/mL)	Articular cavity injection	Once a day for 2 weeks	~5 w (20 d + 2 w)				
20	Jin et al, 2020a	In vivo	Sham OA Exo Exo-mimic NC Exo-miR-9-5p mimic Exo-inhibitor NC Exo-miR-9-5p inhibitor Exo-miR-9-5p inhibitor + liposomes miR-9-5p	-	Intraarticular injection of exosomes	-	7 w	miR-9-5p	syndecan-1	Anti-inflammatory and chondroprotective effects	[22]

(Continued)

Table 3 (Continued).

No.	Author	Disease Model	Groups/Treatments	Concentration/Volume	Treatment/Delivery Route	Frequency	Follow Up	microRNAs	Target Gene	Functional Outcome	Ref.
21	Zhang, Qiet al, 2023	In vitro	Control IL-1 β IL-1 β + BMSC-Exo IL-1 β + dECM-BMSC-Exo For rescue investigation: Control IL-1 β IL-1 β + mimic-NC IL-1 β + miR-3473b mimic	5×10^8 /mL	Exosomes incubated with 10^4 chondrocytes	2 h	-	miR-3473b	PTEN	↓OA by preventing apoptosis, ↑anabolism and migration of chondrocytes	[42]
		In vivo	(1) sham group, n= 5 (2) PBS group, n= 5 (3) BMSC-Exo group, n= 5 (4) dECM-BMSC group, n= 5 For rescue investigation: (1) dECM-BMSC-Exo group, n= 5 (2) dECM-BMSC-Exos + antagomir-NC group, n= 5 (3) dECM-BMSC-Exos + antagomir-3473b group, n= 5	10 μ L of dECM-BMSC-Exos 10^{10} particles/mL	Intra-articular injections	Twice a week for 4 weeks	8 w (4+4 w)			Cartilage regeneration	
22	Dong et al, 2024	In vitro	IL-1 β IL-1 β + BMSCs-CM IL-1 β + BMSCsQUE-CM IL-1 β + BMSCsQUE-CM	Exosome	Incubated with chondrocytes for 24 h	48 h	50 h	miR-124-3p	TRAF6	↓chondrocyte injury	[43]
		In vivo	Control group OA group OA + BMSCs-Exo group OA + BMSCs ^{QUE} -Exo group	200 mL exosome	Exosome in OA patients	Twice a week	-				
23	Lin et al, 2021	In vitro	i) Normal culture ii) IL-1 β iii) IL-1 β + dPSc-derived exosomes iv) IL-1 β + miR-140 enriched exosomes	Exosome (5×10^8 particles/mL)	Incubated with cartilage cells	48 h		miR-140-5p	Col2 α 1 Sox9	↓chondrocyte apoptosis	[44]
		In vivo	Blank, n= 6 OA model, n= 6 Exosome, n= 6 Improved-exosome treatment, n= 6	50 μ L exosomes (5×10^{10} particles/mL)	Knee joint cavity injection	Once a week for 4 weeks	1+4+1 w				

24	Liu et al, 2023	In vitro	IL-1 β IL-1 β + miR-EVs IL-1 β + collagen II-targeting peptides-EVs IL-1 β + collagen II-targeting peptides/miR-EVs Nigericin sodium salt	EVs (5×10^8 particles/mL)	Incubated with chondrocytes	24 h	-	miR-223	NLRP3	↓ cell death ↑ anabolism of chondrocytes ↓ OA severity ↑ miR-223 in the cartilage of exosome-treated knees ↑ chondrocyte homeostasis ↑ protect articular cartilage from damage	[45]
		In vivo	PBS, n= 3 hUC-EVs, n= 3 miR-EVs, n= 3 CTP-EVs, n= 3 CTP/miR-EVs, n= 3 Control, n= 3	50 μ L EVs (10^{10} particles/mL)	Intra-articular injection	Once a week for 4 weeks	5 w				
25	Liu et al, 2022	In vitro	-Blank IL-1 β IL-1 β + Exos (1 μ g/mL) IL-1 β + Exos (2.5 μ g/mL) IL-1 β + Exos (10 μ g/mL) -Blank IL-1 β IL-1 β + hUSC-Exos IL-1 β + hUSC-140-Exos	1, 2.5, or 10 μ g/mL	Incubation	48 h	72 h	miR-140-5p	VEGFA	↓ OA progression by modulating ECM homeostasis and subchondral bone remodeling	[46]
		In vivo	(1) sham, n= 10 (2) OA + PBS, N= 10 (3) OA + hUSC-Exos, n= 10 (4) OA + hUSC-140-Exos, n=10	100 mL (10^{11} particles/mL)	IA injection	Once a week for 4 weeks	8 w				
26	Meng et al, 2023	In vitro	ACs + Exo ^{ADSCs} ACs + TE-Exo ^{ADSCs} ACs + TE + Exo ^{ADSCs} ACs + TE	Exosome (20 μ g/mL)	Exosomes incubated with ACs (5×10^4 cells)	48 h	-	miR-451-5p	KEGG pathway	-maintain the chondrocyte phenotype ↑ cartilage repair	[47]
		In vivo	(1) normal group, n= 5 (2) OA group, n= 5 (3) OA + Exo ^{ADSCs} group, n= 5 (4) OA + TE-Exo ^{ADSCs} group, n= 5 (5) OA + TE + Exo ^{ADSCs} group, n= 5 (6) OA + TE group, n= 5	Exosomes	Intra-articular injection	4 w	8 w				

(Continued)

Table 3 (Continued).

No.	Author	Disease Model	Groups/Treatments	Concentration/Volume	Treatment/Delivery Route	Frequency	Follow Up	microRNAs	Target Gene	Functional Outcome	Ref.
27	Sun et al, 2022	In vitro	Control TGF- β 3 un-TGF β 3- preconditioned EV group TGF β 3-preconditioned EV group	EV (1×10^8 , 2×10^8 particles/ mL)	Incubated with 4×10^4 cells/well	12 h	7 d	miR-455	SOX11/ FOXO axis	\uparrow OA injury and cartilage regeneration	[48]
		In vivo	OA, n= 8 OA + untreated-EVs, n= 8 OA + TGF β 3-treated EVs, n= 8	50 μ L EVs (10^{10} particles/mL)	Intraarticular injection EVs	Once a week for 7 w (5 th -12 th)	24 w				
28	Wu et al, 2019	In vitro	Control IL-1 β IL-1 β + MSC ^{IPFP} -Exo1 IL-1 β + MSC ^{IPFP} -Exo2 IL-1 β + MSC ^{IPFP} -Exo3	1, 5, or 10×10^8 exosome particles/mL	Exosomes incubated with SW1353 cells	24 h	-	miR-100-5p	mTOR	-Protect cartilage from damage \downarrow gait patterns of DMM-induced OA mice \downarrow chondrocyte apoptosis -balancing the anabolic and catabolic processes	[49]
		In vivo	(1) sham group (2) PBS group (3) PBS-Exo ^{IPFP} group (1) PBS + antagomir-NC group (2) PBS-Exo ^{IPFP} + antagomir-NC group (3) PBS-Exo ^{IPFP} + antagomir-100-5p group	-10 μ L exosomes (10^{10} particles/mL) -10 μ L antagomir and then 10 μ L Exos	Intraarticular injection	- twice a week for 4 or 6 weeks - antagomir once a week for 3 weeks and then exo twice a week for 4 weeks	6 w				
29	Ye et al, 2022	In vitro	-Blank IL-1 β IL-1 β + MSC-EV -Blank IL-1 β IL-1 β + MSC-EV-mimic- NC IL-1 β + MSC-EV- miR- 3960 mimic	100 μ g/mL MSCs-EVs	EVs incubated with chondrocytes	36 h	2-4 d	miR-3960	PHLDA2	Altered the SDC1/ Wnt/ β -catenin expression \downarrow chondrocyte injury in OA	[50]
		In vivo	Sham, n= 8 OA, n= 8 OA + MSCs-EVs- agomirNC, n= 8 OA+ MSCs-EVs-miR- 3960 agomir, n= 8	10 μ g/mL	Injection	3 w	7 w				

30	Meng et al, 2023	In vitro	NC IL-1 β miR-429 mimics pcDNA3.1-FE22 miR-429 mimics + pcDNA3.1-FE22	100 μ g/mL	Exosomes incubated with chondrocytes	48 h	-	miR-429	FEZ2	\uparrow chondrocyte proliferation \uparrow autophagy	[51]
		In vivo	Control, n= 10 OA, n= 10 NC, n= 10 miR-429 mimics, n= 10 pcDNA3.1-FE22, n= 10 miR-429 mimics + pcDNA3.1-FE22, n= 10	40 μ g	Knee joints injection	Once a week for 6 weeks	6 w				
31	Zhao et al, 2023	In vitro	Control, n= 6 IL-1 β + PBS, n= 6 IL-1 β + Exo ^{SC} , n= 6 IL-1 β + CAP-Exo ^{SC} , n= 6 IL-1 β + Exo ^{SC} -miR-199, n= 6 IL-1 β + CAP-Exo ^{SC} -miR-199, n= 6	1×10^9 particles/mL	Exosomes incubated with rat synovial cells (1×10^5 cells/well)	24 h	48–72 h	miR-199a-3p	mTOR	\uparrow chondrocyte autophagy \uparrow anabolism \downarrow catabolism \downarrow pathological severity	[52]
		In vivo	(1) sham (2) PBS + antagomir-NC (3) PBS-ExosSC + antagomirNC group (4) PBS-ExosSC + antagomir-199-3p group (1) sham group (2) OA+PBS group (3) OA+ExosSC group, (4) OA+ExosSC/mir-199a-3p group (5) OA+CAP-ExosSC/ mir-199a-3p group	- pre-injected with 50 μ L antagomir-NC orantagomir-199-3p and then 50 μ L PBS or 50 μ L ExosSC (2×10^{10} particles/mL) -10 μ L ExosSC (1×10^{10} particles/mL)	Intraarticular injection	-re-injection for 3 weeks twice a week, 2 week after surgery, then multiple injections of ExosSConce a week during the following 6 weeks. -Multiple injections 6 week after operation	-11 w -10 w				
32	Wang et al, 2020	In vitro	Sh-NC Sh-PTTG1 Sh-PTTG1 + EVs (inhibitor-NC) Sh-PTTG1 + EVs (miR-31 inhibitor)	Treated with EVs for 24 h	EVs incubated with chondrocytes (2.5×10^4 cells/well)	24 h	72 h	miR-31	KDM2A	\uparrow proliferation and migration of chondrocytes -activated the E2F1/ PTTG1 axis: \uparrow chondrocyte proliferation \downarrow expression of inflammatory factors IL-1 β , IL-6, and TNF- α \downarrow occurrence of knee OA	[53]
		In vivo	(1) the sham group, n= 10 (2) the OA group, n= 10 (3) the OA + SMSC-EV group, n= 10 (4) the OA + EV (miR-31 mimic) group, n= 10	EV (5 μ L particles/mL)	Intraarticular injection of EVs	Every 3 days for 4 weeks	12 w				

(Continued)

Table 3 (Continued).

No.	Author	Disease Model	Groups/Treatments	Concentration/Volume	Treatment/Delivery Route	Frequency	Follow Up	microRNAs	Target Gene	Functional Outcome	Ref.
33	Lou et al, 2023	In vitro	Control IL-1 β IL-1 β + BMSC-Exo IL-1 β + F-BMSC-Exo	Exosomes (0, 1, 5, 10 μ g/mL)	Exosomes incubated with rat chondrocytes	12 h	72 h	miR-146b-5p	TRAF6	↓cartilage inflammation, degrading the extracellular matrix, and activating autophagy -inhibiting the PI3K/AKT/mTOR pathway	[54]
		In vivo	-Sham group, n= 6 OA group, n= 6 OA +MSCs-Exo group, n= 6 OA+ F-MSCs-Exo group, n= 6 -the sham group, n= 6 OA group, n= 6 OA+F-MSCs-Exo group, n= 6 OA +F-MSCs-Exo +Antagomir-146b-5p group, n= 6	Exosomes (10 μ L/week) (+ Antagomir-146b-5p (5 nmol))	Intraarticular injection of exosomes	Once a week for 4 weeks	4+4 w				
34	Qiu et al, 2020	In vitro	1. Control 2. IL-1 β 3. IL-1 β + EXO 4. IL-1 β + EXO- curcumin	Exosomes (80 μ g/mL)	Incubated with chondrocytes		48 h	miR-124 miR-143	NF- κ B ROCK1	Regulates the miR-124/NF- κ B and miR-143/ROCK1/TLR9 pathways	[55]
		In vivo	Sham group OA group OA + EXO group OA + EXO- curcumin group	Exosomes	Mice treated with exosomes						
35	Liu et al, 2018	In vitro	-Control IL-1 β PBS MSC ^{vector} -Exos MSC KLF ³ -AS ¹ -Exos - mimic NC + MSC ^{vector} -Exos miR-206 mimic + MSC ^{vector} -Exos mimic NC + MSC KLF ³ -AS ¹ -Exos miR-206 mimic + MSC KLF ³ -AS ¹ -Exos -normal group OA group MSC-Exos group PBS group	Exosome (10 μ g/mL)	2 \times -Exosomes incubated with mice chondrocytes -Adult SPF 9-week-old C57BL/6 mice	24 h	-	lncRNA-KLF3-AS1	miR-206/ GIT1 axis	↑cartilage repair: ↑chondrocyte proliferation and ↓apoptosis	[56]

36	Xia et al, 2021	In vitro	Mimics NC Mimics miR-125a-5p Inhibitor NC Inhibitor miR-125a-5p Mimics miR-125a-5p- E2F2	-	Exosomes incubated with chondrocytes	6 h	-	miR-125a-5p	E2F2	↓endogenous expressions of E2F2 and MMP-1 ↓expressions of collagen II, aggrecan, and Sox9 ↑chondrocyte migration ↓cartilage degeneration	[57]
		In vivo	Normal, n= 12 OA, n= 12 OA + exosomes, n= 12 OA + ExosmiR -125a-5p, n= 12	100 μ L of 10^{11} particles/mL	Joint injection of exosomes	-	-				
37	Tao et al, 2021	In vitro	Exo-miR-NC exo-miR-361-5p exo-miR-361-5p + vector exo-miR-361-5p + DDX20	2 μ g exosomes	2 μ g exosomes incubated with chondrocytes for 48 h	48 h	-	miRNA-361-5p	DDX20 and NF- κ B signaling pathway	↓inflammation of chondrocytes -protective effect against OA	[58]
		In vivo	Sham, exo-miR-NC group, n= 15 Sham, exo-miR-361-5p group, n = 15 OA, exo-miR-NC group, n= 15 OA, exo-miR-361-5p group, n = 15	250 ng/5 μ L	Joint injection	7 d	8 w				
38	Chen et al, 2020	In vitro	Agomir-NC miR-136-5p agomir antagomir-NC miR-136-5p antagomir miR-136-5p agomir + ELF-3	-	Exosomes incubated with chondrocytes for	6 h	-	miR-136-5p	ELF3	↓expressions of ELF3 and MMP-1 ↑levels of collagen II, aggrecan, and SOX9 ↑migration of chondrocytes ↓cartilage degeneration	[13]
		In vivo	Normal, n= 12 Post-traumatic oleonic acid (OA), n= 12 Post-traumatic OA + exosomes, n= 12 Post-traumatic OA + ExosmiR-136-5p, n= 12	100 μ L of exosomes (10^{11} particles/mL)	Joint injection	1 h	1 h				
39	Lu et al, 2021	In vitro	GW, EVs, EVs-negative control (NC), or EVs-inhibitor	30 μ L EVs (10^{11} particles/mL)	Incubated with SW1353 cells	6 h	24 h	miR-26a-5p	PTEN	↓apoptosis and inflammation ↓cartilage damage of OA	[59]
		In vivo	Sham group, n=6 OA group, n= 6 OA + GW group, n= 6 OA + EVs group, n= 6 OA + EVs-NC group, n= 6 OA + EVs-inhibitor group n=6	30 μ L EVs (10^{11} particles/mL)	Intraarticular injection	Days 7, 14, and 21	28 d				

(Continued)

Table 3 (Continued).

No.	Author	Disease Model	Groups/Treatments	Concentration/Volume	Treatment/Delivery Route	Frequency	Follow Up	microRNAs	Target Gene	Functional Outcome	Ref.
40	Tao et al, 2017	In vitro	SMSC-140-Exos-0 SMSC-140-Exos-1 SMSC-140-Exos-5 SMSC-140-Exos-10 empty vector + control empty vector + SMSC-Exos-10 shYAP #1 + control shYAP #2 + control shYAP #1 + SMSC-Exos-10 shYAP #2 + SMSC-Exos-10	-0, 1, 5, or 10×10^{11} particles/mL -10×10^{11} particles/mL	Exosomes incubated with chondrocytes (2×10^4 cells/well)	24 h	12 h	miR-140-5p	SOX-9 YAP	↓progression of early-stage OA ↓knee joint cartilage damage	[24]
		In vivo	(1) Normal group, n = 10 (2) OA group, n = 10 (3) OA+SMSC-Exo group, n = 10 (4) OA+SMSC-140-Exo group, n = 10	100 μ L exosomes (10^{11} particles/mL)	Intraarticular injection	First day of every w from the 5 th -8 th w after surgery	12 w				
41	Kong et al, 2023	In vivo	DMM group, n=3 Low-dose SMSC-Exo group, n=3 Medium-dose SMSC-Exo group, n=3 High-dose SMSC-Exo group, n=3	Exosomes (50 μ g, 100 μ g or 200 μ g in 50 μ L normal saline)	Articular cavity injection	4 w	8 w	miR-320c	ADAM19-dependent Wnt signalling	↓ECM degradation and chondrocyte apoptosis; ↑cartilage damage repair	[60]

Abbreviations: ADAMTS9, ADAM metalloproteinase with thrombospondin type 1 motif 9; ADSCs, adipose-derived stromal cells; ATG4A, autophagy related 4A; AVN, Avascular Necrosis or Osteonecrosis; BMSCs, Bone marrow mesenchymal stem cells; CDH11, Cadherin-11; CIA, Collagen-induced arthritis; CXCL9, Chemokine ligand 9; DDX20, DEAD-Box Helicase 20; E2F2, E2F Transcription Factor 2; ELF3, E74 Like ETS Transcription Factor 3; GIT1, G-protein-coupled receptor kinase interacting protein-1; FLS, Fibroblast-like synovocyte; GREM1, Gremlin 1; IGF1R, Insulin-like growth factor1 receptor; hUC-MSCs, Human umbilical cord-derived mesenchymal stem cells; KLF, Krüppel-like factor; MAPK6, Mitogen-activated protein kinase 6; MDM2, Mouse double minute 2 homolog; MMP13, Matrix Metalloproteinase 13; mTOR, mechanistic Target of Rapamycin; MSC, Mesenchymal stem cell; NF- κ B, Nuclear Factor Kappa B; NOX4, NADPH Oxidase 4; OA, Osteoarthritis; PTGS2, prostaglandin-endoperoxide synthase 2; RA, Rheumatoid arthritis; ROS, Reactive oxygen species; SGK1, Serum/Glucocorticoid Regulated Kinase 1; hSMSCs, human synovial mesenchymal stem cells; TET1, Ten-eleven translocation methylcytosine dioxygenase 1; TGF- β , Transforming growth factor Beta; TNF- α , Tumor necrosis factor alpha.

Table 4 Preclinical Studies Applied MSC-Derived Exosomal microRNAs for Rheumatoid Arthritis-Additional Information

No.	Author	Disease Model	Groups/Treatments	Concentration/ Volume	Treatment/Delivery Route	Frequency	Follow Up	microRNAs	Target Gene	Functional Outcome	Ref.
1	Tavasolian et al	In vitro	PHA miR-146 miR-155 Exosome Scramble NC	Exosomes (10 mg/mL)	ExoSome co-culture	At d 0	72-120 h	miR-146a	TRAF6	↑Treg cell populations ↑anti-inflammatory factors	[15]
2	Meng et al 2020	In vitro	1. MH7A cells + BM- MSC-124a-EV 2. MH7A cells + BM- MSC-EV 3. MH7A cells + Control	20 µg/mL	MH7A cells treated with EVs	At d 0	48 h	miR-124a	Caspase 3,9 Bax, Bid, Bim, Bcl-2	↑apoptosis ↓proliferation and migration of FLS	[61]
3	Su et al, 2021	In vitro	1. MSC-HAND2-ASI- Exos 2. vector-Exos 3. Control	50 µg/mL	Exosomes incubated with 4 × 10 ⁵ MH7A cell/well	At d 0	48 h	lncRNA HAND2-ASI	↓miR-143-3p/ ↑TNFAIP3/ ↓NF-κB pathway	↓proliferation, invasion, migration, and inflammation ↑apoptosis	[62]
4	Chen et al, 2018	In vitro	1. MSC-Exo-150 2. MSC-Exo-67 (control)	10 µg/mL	RA-FLS and HUVEC cells treated with MSC-Exo	At d 0	48 h	miRNA-150-5p	MMP14 and VEGF	↓migration and invasion in RA FLS ↓tube formation in HUVECs	[63]
		In vivo	1. MSC-Exo-150 2. MSC-Exo-67 (control) 3. PBS	100 µL PBS containing 50 µg exosomes	IP injection	Twice a week	35 d			↓synovocyte hyperplasia and angiogenesis ↓hind paw thickness ↓clinical arthritis scores	

(Continued)

Table 4 (Continued).

No.	Author	Disease Model	Groups/Treatments	Concentration/Volume	Treatment/Delivery Route	Frequency	Follow Up	microRNAs	Target Gene	Functional Outcome	Ref.
5	Huang et al, 2022	In vitro	-NC LPS + ATP LPS + ATP + exosomes -Exo-miRNA-NC inhibitor Exo-miRNA-223 inhibitor Exo-miRNA-223 inhibitor + si-NC Exo-miRNA-223 inhibitor + si-NLRP3	Cultured medium (10 µM)	Co-cultured	24 h	-	miR-223	NLRP3	Anti-inflammatory effects	[64]
		In vivo	Control, n= 8 Arthritis model (AA), n= 8 AA + DXMS, n= 8 AA + Exosomes, n= 8	Exosomes	-	Once a day for 28 days	28 d				
6	Lin et al, 2023	In vitro	Control Mimic NC Anti-miR-NC Anti-miR-433-3p	sEV (100 µg)	Incubated with HDMECs (5 × 10 ⁵ cells/mL)	48 h	-	miR-433-3p	FOXO1	↓synovial angiogenesis	[65]
		In vivo	Control group Erastin-sEV SMSCs-sEV miR-433-3p-SMSCs-sEV Erastin-sEV + miR-433-3pSMSCs-sEV	sEV (80 µg)	Intraarticular injection	Twice a week for 17 days	39 d				

7	Huang et al, 2022	In vitro	1. RASFs (control) 2. RASFs+ HU-MSCs-exo 3. RASFs+ HU-MSCs-transfected with miR-140-3p mimic NC 4. RASFs+ HU-MSCs-transfected with miR-140 mimic-exo 5. RASFs transfected with miR-140 mimic NC 6. RASFs transfected with miR-140 mimic 7. RASFs transfected with miR-140-mimic and SGK1 vector	-	Exosomes incubated with RASFs (1×10^6 cells/mL)	At d 0	3 d	microRNA-140-3p	SGK1	↓RASF growth	[66]
		In vivo	1. Sham 2. CIA 3. Exo 4. miR-140 mimic-exo 5. miR-140 inhibitor-exo 6. mimic-NC-exo 7. inhibitor-NC-exo	100 µL exosomes or miRNA-transfected exosomes (1 µg/µL)	Injected with exosomes or miRNA-transfected exosomes every week (1 µg/µL)	Every week	42 d			↓joint injury ↓chondrocyte ↓apoptosis ↓ fibrosis ↓ oxidative stress ↓inflammation	
8	Mi et al, 2024	In vitro	1. Control, 2. hUC-MSC-Exos, 3. hUC-MSC-ExosmiR-451a	150 µg/mL	Exosomes incubated with RA SF (3×10^3 cells/well)	At d 0	48h	miR-451a	ATF2	↓proliferation, migration, and invasion of hSFs	[67]
		In vivo	1. WT 2. CIA 3. hUC-MSC-ExosmiR-451a 4. hUC-MSC-Exos	-	Transplant of ExosmiR-451a to ankle joint cavities rats	At 16 and 24 day	45 d			↓Joint inflammation	
9	Meng and Qiu, 2020	In vitro	1. PBS 2. Exo 3. Exo- mimic-NC 4. Exo- miR-320a mimic	100 µg	Exosomes incubated with RA-FLSs (5×10^4 cells/well)	At d 0	72h	miR-320a	CXCL9	↓RA-FLS activation, migration, and invasion	[68]
		In vivo	1. Normal 2. CIA 3. CIA-PBS 4. CIA-Exo-miR-NC 5. CIA- Exo- miR-320a mimic	100 µg/day	Injected into the tail vein of mice	20 d	21d			↓arthritis and bone damage	

(Continued)

Table 4 (Continued).

No.	Author	Disease Model	Groups/Treatments	Concentration/Volume	Treatment/Delivery Route	Frequency	Follow Up	microRNAs	Target Gene	Functional Outcome	Ref.
10	Ma et al, 2022	In vitro	1. BM-MSCs-exosomes 2. Chondrogenic BM-MSCs secreted exosomes	50 nM	Incubated with FLSs (4×10 ⁵ cells/well)	At d 0	48 h	miR-205-5p	MDM2	↓MAPK and NF-κB pathways	[69]
		In vivo	1. Normal 2. CIA 3. CIA+exo-miR-NC 4. CIA+exo-miR-205-5p	200 µL	Intradermal injection of exosomes to mice tail	Twice a week	14 d			↓inflammation score and response ↓joint destruction	
11	Li et al, 2021	In vitro	1. Control 2. Exo (inhibitor NC) 3. Exo (miR-21 inhibitor)	-	EVs incubated with mFLS cells	At h 0	6 h	miR-21	TET1	↓KLF4	[70]
		In vivo	1. Normal, 2. RA, 3. RA +Exo (inhibitor NC), 4. RA+Exo (miR-21 inhibitor)	50 mg	Injection of EVs to mice	Twice a week	16 d			Improve RA progression in mice	
12	Zhang et al, 2023	In vitro	1. Control 2. Mimic NC 3. EVs-miR-378a-5p mimic 4. Inhibitor NC 5. EVs-miR-378a-5p inhibitor	140 µg/mL	EVs added to HSMECs	20 h	48 h	miR-378a-5p	Blocks IRF1/STAT1 axis	↑angiogenesis, migration, and proliferation of HSMECs	[71]
		In vitro	1. Sham 2. CIA 3. CIA + EVs-miR-378a-5p mimic 4. CIA+ dexametomidine 5. CIA + EVs-miR-378a-5p mimic + dexametomidine	140 µg/mL	EVs 0.5 mL/mouse/day intraperitoneal injection	6 d	20 d			↑synovial vascular remodeling and histopathology	

13	Wu et al, 2021	In vitro	1. EVs 2. EVs- miR-negative control 3. EVs- miR-34a inhibitor	75 µg/mL	EVs treated with FLSs	At h 0	36 h	miR-34a	↓Cyclin I ↑ATM/ATR/ p53 axis	↓proliferation ↑apoptosis ↓Inflammation	[72]
		In vivo	1. EVs group 2. Normal 3. EVs- miR-negative control 4. EVs- miR-34a inhibitor	75 µg/mL	Tail vein injection of EVs	At d 1	4 weeks				
14	Wu et al, 2022	In vivo	-	-	-	-	-	miR-23-5p	RAC2	↓Inflammation	[73]

Abbreviations: ADAMTS9, ADAM metallopeptidase with thrombospondin type 1 motif 9; AVN, Avascular Necrosis or Osteonecrosis; BMSCs, Bone mesenchymal stem cells; CDH11, Cadherin-11; CXCL9, Chemokine ligand 9; DDX20, DEAD-Box Helicase 20; E2F2, E2F Transcription Factor 2; ELF3, E74 Like ETS Transcription Factor 3; FLS, Fibroblast-like synoviocyte; GREM1, Gremlin 1; IGF1R, Insulin-like growth factor I receptor; hUC-MSCs, Human umbilical cord-derived mesenchymal stem cells; KLF, Krüppel-like factor; MAPK6, Mitogen-activated protein kinase 6; MDM2, Mouse double minute 2 homolog; MMP13, Matrix Metallopeptidase 13; mTOR, mechanistic Target of Rapamycin; MSC, Mesenchymal stem cell; NF-κB, Nuclear Factor Kappa B; NOX4, NADPH Oxidase 4; OA, Osteoarthritis; PTGS2, prostaglandin-endoperoxide synthase 2; RA, Rheumatoid arthritis; ROS, Reactive oxygen species; SGK1, Serum/Glucocorticoid Regulated Kinase 1; hSMSCs, human synovial mesenchymal stem cells; TET1, Ten-eleven translocation methylcytosine dioxygenase 1; TGF-β, Transforming growth factor Beta; TNF-α, Tumor necrosis factor alpha; ADSCs, adipose-derived stromal cells; ATG4A, autophagy related 4A; GIT1, G-protein-coupled receptor kinase interacting protein-1.

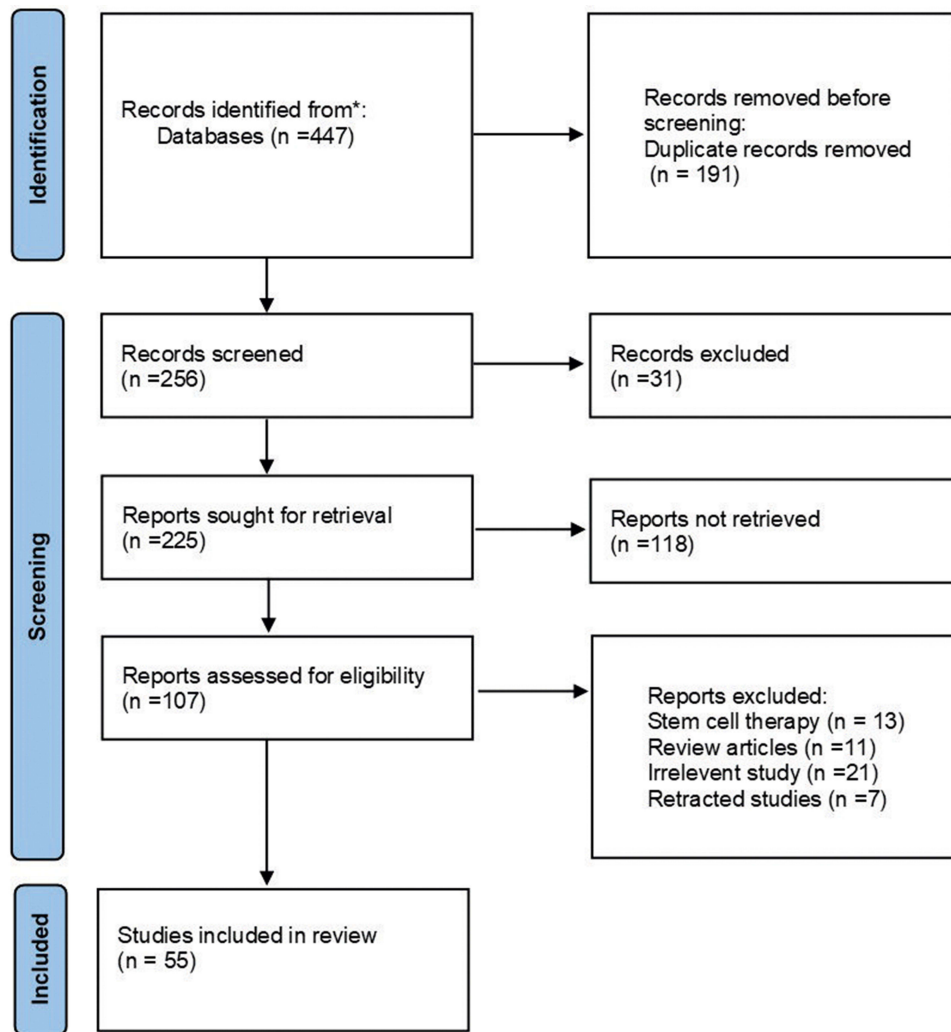


Figure 1 PRISMA flow diagram of the literature search and selection process. *Web of Sciences, PubMed, Scopus, and Google Scholar.

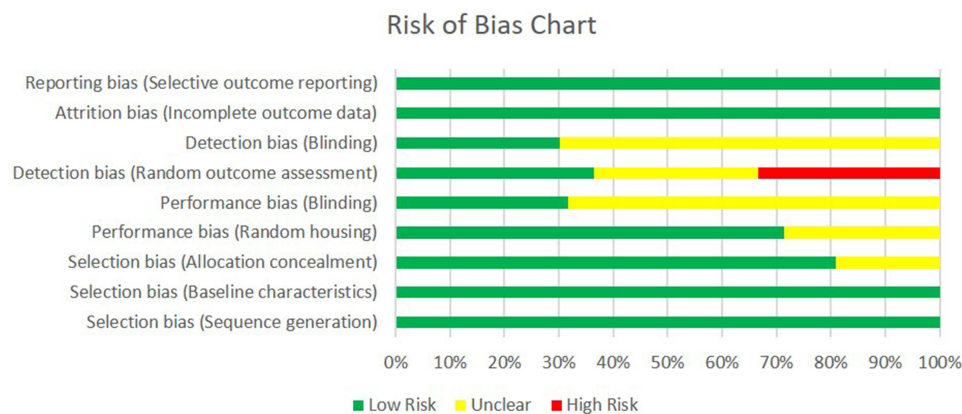


Figure 2 Risk of bias chart.

MSC Sources and Pre-Conditioning Methods to Obtain Engineered Exosomes in RA

Fourteen studies in total were conducted to assess the impact of MSC-derived EVs on the treatment of RA, with a detailed description of their characteristics found in Table 2. Among these studies, three were in vitro studies,^{15,61,62} ten

utilized a combination of methodologies (in vitro+ in vivo,^{63–72} and one was in vivo study.⁷³ The MH7A cell line was selected as a dependable in vitro model for replicating Fibroblast-like synoviocytes from RA patients. Various animal models, including mice (n=6)^{63,65,68–71,73} and rats (n=4)^{64,66,67,72} were employed. Collagen-induced arthritis (CIA) was the most widely used model for studying RA.

The included RA studies showcased a range of MSC types. The most commonly utilized MSCs were BM-MSCs (n=10),^{61–64,68–73} followed by umbilical cord MSCs (UC-MSCs),^{66,67} AD-MSCs,¹⁵ and synovial MSCs (S-MSCs).⁶⁵ In most studies, MSCs were transfected with specific miRNA mimics or inhibitors to generate engineered exosomes, [Table 2](#).

Isolation and Characterization of Exosomes

The ultracentrifugation technique was predominantly used to extract exosomes from MSCs, although some studies opted for different exosome extraction kits like Invitrogen,⁶¹ ExoQuick-TC,^{21,34,35,40,42,49,58,63,66} or HiefTM⁶⁰ kits. Transmission electron microscopy (TEM), Western blotting, and nanoparticle tracking analysis were utilized to examine the structure, presence of exosome-related markers, and size distribution, respectively. The extracted MSC exosomes typically exhibited a bilayer lipid membrane structure under the TEM, showcased a size distribution ranging from around 30 to 200 nm, and harbored exosome-related markers, including endosomal proteins ALIX and TSG101, tetraspanins CD81, CD63, and CD9 ([Table 1](#) and [Table 2](#)).

Delivery Routes and Treatment Frequency for miRNAs Derived from MSC-Exosomes

A wide range of exosome concentrations, ranging from 20–200 µg/µL, were utilized across the studies. The intraperitoneal, intraarticular, intradermal, and tail vein routes were employed for the administration of exosomes. Additionally, in one investigation, exosomes were transplanted into ankle joint cavities.⁶⁷ Furthermore, in certain instances, the concentration of exosomes was unspecified, with only the total quantity of injected exosomes being referenced. The exact dosage and route of administration were not elucidated in some studies. In animal experiments, the duration of observation varied between 14 and 45 days after treatment ([Table 3](#) and [Table 4](#)).

Outcomes/Improvement

Generally, the results indicated that the administration of MSC-exosomal miRNAs had a positive impact on cartilage regeneration and cartilage repair in the defect area. Moreover, studies illustrated that MSC-secretome therapy of OA patients could reduce apoptosis, pyroptosis of cartilage, ROS production, pathological severity, inflammation, and degrading of the extracellular matrix in the chondrocytes. This therapy could also increase the production of cartilage, homeostasis, chondrocyte proliferation, differentiation, migration, autophagy of chondrocytes ([Table 3](#)). Based on the obtained results of experimental RA studies, it was observed that MSC-secretome decreased proliferation, invasion, migration, inflammation, hyperplasia, angiogenesis, and arthritis scores of synoviocyte cells ([Table 4](#)).

Quality and Risk of Bias Assessment

Considerable heterogeneity was observed among the included studies in several key methodological aspects. Various sources of MSCs, including bone marrow, adipose tissue, umbilical cord, and synovium, potentially influence the composition and biological potency of secreted exosomes. Exosome isolation and characterization techniques were not standardized; studies employed differential ultracentrifugation, precipitation kits, size-exclusion chromatography, or microfiltration, leading to variations in particle purity and yield. Animal models ranged from chemically induced to surgically created OA and RA models, with differences in disease severity and duration that may affect treatment response. Likewise, the dosage and administration routes of exosomes (intra-articular injection, intravenous infusion, or local tissue delivery) varied substantially, limiting direct comparability across studies. These methodological inconsistencies contribute to variability in reported outcomes and complicate the identification of dose-response relationships or optimal treatment regimens. As a result, while consistent beneficial effects of MSC-derived exosomal miRNAs were reported, caution is warranted when generalizing these findings across experimental systems or translating them to clinical settings.

In the included studies, the main risk of bias was the lack of clear information about the blinding and randomization process in the performance and detection phase (Figure 2). Blinding of investigators and outcome assessors was rarely implemented, particularly for histological scoring or behavioral analyses, which represent a risk of detection bias. Only a few studies provided sample size justification, technical replicates, or independent repeat experiments, which limit reproducibility.

Discussion

MSC-Derived EVs in Osteoarthritis

The regenerative and immunomodulatory role of MSC-EVs in OA has attracted the attention of many researchers. Accordingly, several studies have revealed the significant therapeutic role of MSC-EVs in OA by suppressing degenerative pathways and specific pro-inflammatory elements and improving the migration and proliferation of chondrocytes.^{74,75} By inhibiting apoptosis, pyroptosis, and ferroptosis, they can inhibit chondrocyte death. These cargo-carrying components are also capable of improving chondrogenesis and cellular differentiation and stimulating autophagy to preserve chondrocyte integrity, which is directly correlated with chondrocyte regeneration and efficient metabolism (reviewed by Wu et al).⁷⁶ MSC-derived exosomes have also been demonstrated to reduce the anabolic capacity of chondrocytes.²⁴ It is also reported that overexpressing miR-140 in urine-derived MSCs could ease knee OA in a rat model by downregulation of endothelial growth factor A.⁴⁶

Vonk et al revealed that MSC-EVs are capable of blocking NF- κ B signaling by suppressing phosphorylation of I κ B α , leading to downregulation of TNF- α (tumor necrosis factor-alpha)-derived COX2 activity, cytokine-mediated responses, and collagenase function.⁷⁴ Additionally, MSC-EVs are involved in the upregulation of SOX9 and WNT7A expression. In vitro synthesis of proteoglycan and type II collagen is enhanced by MSC-EVs.⁷⁴ Induction of OA-like chondrocytes with MSC-EVs is associated with an elevated level of chondrogenesis (type II collagen and aggrecan) indicators, and reduced level of catabolic (MMP-13), and inflammatory (iNOS) factors.⁷⁵ In another study, the MSC-EV treatment of OA in mouse models was associated with decreased cartilage and bone degradation.⁷⁷ Ruiz et al also demonstrated that the therapeutic impact of MSC-EVs in a mouse model of OA was partially regulated by TGF- β 1.⁷⁸ Further studies have shown that miR-206, miR-212-5p, and miR-136-5p present in the MSCs-derived exosomes enhance chondrocyte migration, prevent chondrocyte degeneration, and decline inflammatory responses in OA models by targeting E74-like factor 3 (ELF3).^{13,37,40} ELF3 is an inflammatory mediator activated by pro-inflammatory cytokines in chondrocytes, which elevates the expression level of inflammatory and catabolic pathways, involved in cartilage degradation in OR. EVs obtained from MSCs with overexpressed circHIPK3 (EVs-circHIPK3) are associated with promotion of chondrocyte migration and proliferation and suppression of chondrocyte apoptosis induced by the miR-124/MYH9 pathway.⁷⁹

Exosomal-miRNAs Derived from BM-MSCs in Osteoarthritis

Various studies have demonstrated that BM-MSC-exosomal miRs have therapeutic effects on OA by attenuating inflammatory-mediated cartilage injury, promoting proliferation and migration and suppressing apoptosis of chondrocytes in OA joints. BM-MSC-exosomal miRNAs, such as miR-136-5p,¹³ miR-6515-5p³² prevent the apoptosis and senescence of chondrocytes. Surprisingly, one study found that in HIF-1 α -induced BM-MSC-EVs, hypoxia results in overexpression of miR-216a, leading to migration and proliferation, and a drop in apoptosis of chondrocytes in OA (through suppression of the JAK2 and STAT3 signaling pathway).³⁰ Hu et al indicated that under hypoxia, the overexpression of miR-140-3p in exosomes derived from BM-MSCs could significantly increase migration, survival rate, and chondrogenic differentiation in inflammatory chondrocytes and consequently delay OA pathogenesis and protect joints.²⁶ On the whole, BM-derived exosomal miR-9-5p, miR-92a-3p, miR-127-3p, miR-210, miR-135b, miR-326, miR-206, miR-361-5p, miR-347-3b, miR-124-3p, miR-146-5p, miR-124, miR-143, miR-455, miR-125a-5p, and miR-3960 by decreasing inflammation, cell death, and inducing the proliferation and migration of chondrocytes represent therapeutic effects on OA.^{21,22,25,26,30,33,35,38,40,42,43,48,50,54,55,57,58,72}

Exosomal-miRNAs Derived from AD-MSC in Osteoarthritis

AD-MSCs-exosomes are uptaken by chondrocytes⁸⁰ and reduce the catabolic state in these cells,⁸¹ making them potential therapeutic candidates for OA. Experiments on animal models of OA have demonstrated that AD-MSCs-EVs are capable of suppressing inflammation by inhibiting M1 macrophage infiltration to joint synovium and protecting against cartilage degeneration in both subacute and chronic animal models of OA.⁸¹ Early stages of OA are associated with inflammatory-mediated catabolism in joint cartilage by proteases that can be reversed by IR injection of AD-MSCs-EVs. They repress the progression of OA, which is mediated by enhancing collagen II synthesis and downregulation of ADAMTS-5, MMP-13, MMP-3, and MMP-1 in the presence of IL-1 β .⁸¹ AD-MSCs-EVs are also capable of neutralizing the IL-1 β -mediated inflammatory responses by inhibiting p65 in the NF- κ B signaling pathway.⁸² Exosomes derived from pretreated AD-MSCs with Tropolastin could promote cartilage repair in animal models of OA and preserve the chondrocyte phenotype in cell culture through miR-451-5p.⁴⁷

Comparing the effect of AD-MSCs-EVs on chondrocytes and synovial cells showed that the uptake of exosomes by synovial cells happens quicker with greater intensity, which leads to a better modulation of secretion of interleukins, catabolic, angiogenesis, and pain factors in synovial cells. Moreover, the absorption of exosomes by synovial cells is associated with the release of secondary vesicles from these cells that improve the biological function of exosomes via positive feedback.⁸²

IPFP-MSCs are a type of AD-MSCs that have already been demonstrated to contribute to chondrogenesis in OA.⁸³ It is now being proposed that a part of this impact may be due to IPFP-MSCs-derived exosomes. In vitro studies have shown that these miR-100-5p-enriched exosomes contribute to the suppression of apoptosis and improvement of matrix synthesis by inhibiting the mTOR autophagy pathway, leading to reduced cartilage degradation and gait abnormalities in DMM-induced OA models.⁴⁹ Other in vitro investigations have proposed that IPFP-MSC-EVs improve chondrogenesis in periosteal cells by enhancing the miR-145 and miR-221 expression and downregulating the synthesis of pro-inflammatory cytokines.⁸⁴ Overexpressed exosomal miR-199a-3p in chondrocytes is shown to target the 3' UTR region of mTOR mRNA and reduce protein levels of mTOR and its downstream factors along with increasing chondrocyte autophagy, enhance anabolism and suppressing catabolism for the OA treatment.⁵²

Exosomal-miRNAs Derived from Umbilical Cord-MSC in Osteoarthritis

UC-MSCs have the advantages of easy access, minor immune rejection, and significant immunity-regulating effects, which make them a promising treatment for OA. Researchers have found that UC-MSCs-exosomes can prevent apoptosis and ROS production. This may be due to inhibition of NOX4 expression by exosomal miR-100-5p.³⁶ miR-223, which directly interacts with the 3' untranslated region of NLRP3 mRNA, has been identified as a pivotal miR that facilitates the beneficial effects of hUC-EVs in the realms of inflammation suppression and cartilage protection. The dual-engineered EVs demonstrated a maximal capacity for inhibiting chondrocyte pyroptosis and NLRP3 inflammasome activation, thereby yielding remarkable outcomes in the therapeutic management of osteoarthritis.⁴⁵

Exosomal-miRNAs Derived from Synovial-MSC in Osteoarthritis

Liu et al reviewed the potential roles of EVs and their combination with biomaterials in OA pathogenesis, diagnosis, and therapy.⁸⁵ Tao et al indicated that miR-140-5p overexpression could block Wnt-5a and -5b signaling pathway that activates YAP and decreases ECM secretion.²⁴ In rat OA models induced by cold-water stimulation, OA chondrocytes were treated with overexpressed miR-155-5p S-MSCs-exosomes. This therapy could enhance the release of ECM, cellular proliferation and migration and diminish cell apoptosis, effectively prohibiting OA from happening.²³ Moreover, it is indicated that overexpressed miR-155-5p S-MSCs-exosomes could target Runx2 and increase ECM secretion, causing increased cartilage regeneration and decreased OA.²³ Wang et al found that overexpressed miR-31 S-MSCs-EVs could mitigate OA by regulating the KDM2A (lysine demethylase 2A)/E2F1/PTTG1 (pituitary tumor-transforming gene 1) pathway.⁵³ It is also reported that S-MSC-exosomal miR-129 could inhibit the IL-1 β -mediated OA by preventing the release of high mobility group protein -1 (HMGB1).²⁷ Kong et al found that S-MSC-exosomal miR-320c prevents chondrocyte apoptosis and ECM degradation by targeting ADAM19-dependent Wnt signaling to assist

cartilage injury repair in OA rats.⁶⁰ Modified exosomes derived from miR-155-overexpressing S-MSCs by reducing apoptosis and modulating ECM in chondrocytes could prevent OA.²³

MSC-Derived EVs in Rheumatoid Arthritis

BM-MSC-EVs, like their parental cells, have promising therapeutic effects against RA.⁸⁶ The BM-MSC-Exosomal RNAs are absorbed by fibroblast-like synovial cells (FLSs) and prevent inflammation during RA treatment. It is found that RA-FLS inflammation and proliferation were decreased by miR-34 in BM-MSC-EVs through the prevention of cyclin I/ATM/ATR/p53 signaling pathway.⁷² Exosomal miR-320 derived from BM-MSC significantly downregulates the chemokine ligand CXCL9 and prevents the invasion, migration, and activation of RA-FLSs.⁶⁸ FLS regulation and angiogenesis inhibition occur by exosomal miR-150-5p, resulting in downregulated levels of MMP14 and VEGF.⁶³ It is also reported that the combination of HAND2-AS1 (a long non-coding RNA) with BM-MSC-EVs could suppress RA progression by inhibiting the RA-FLSs tumor-like behavior through the miR-143/TNFAIP3/NF- κ B pathway.⁶² MiRNA-223 in BM-MSCs-exosomes regulates the activation of inflammasomes by targeting NLRP3 in macrophages of RA models.⁶⁶ Generally, BM-MSC-EVs are shown to protect against RA but more research is needed to clarify their anti-inflammatory mechanisms and immune regulatory properties in RA.^{61–64,66,68–73}

Yin et al indicated that miR-99 b-3p in exosomes of subcutaneous AD-MSCs could regulate the cartilage ECM degradation via targeting ADAMTS4 during OA treatment.⁸⁷ Li et al revealed that human AD-MSC-exosomal miR-376c-3p could inhibit the Wnt- β -catenin pathway and then alleviate synovial fibrosis and chondrocyte degradation in an OA animal model.⁶⁵ Altogether, the results showed that AD-MSCs-EVs could be a novel proper alternative RA treatment.

Upregulated exosomal miR-140 in UC-MSCs reduced joint damage of RA rats through targeting serum- and glucocorticoid-inducible kinase 1.⁶⁶

Across MSC sources, a convergent set of intracellular signaling pathways emerges as the principal targets of exosomal miRNAs in OA and RA. Many of the most frequently reported miRNAs (such as miR-140-5p, miR-146a, miR-21, miR-92a-3p, and miR-100-5p) modulate key regulatory cascades that collectively govern inflammation, cartilage matrix turnover, and cell survival. NF- κ B signaling is the most consistently inhibited pathway, underpinning the anti-inflammatory and immunosuppressive effects observed across studies. Several miRNAs (eg, miR-146a, miR-21, miR-320a) attenuate NF- κ B activity by targeting upstream mediators such as TRAF6 and IRAK1, thereby reducing cytokine production and synovial inflammation. Wnt/ β -catenin and PI3K/AKT/mTOR pathways are recurrently modulated to promote anabolic and anti-apoptotic effects in chondrocytes: miR-92a-3p and miR-100-5p restore cartilage matrix synthesis via activation of SOX9 and COL2A1, while inhibition of mTOR signaling by miR-100-5p enhances autophagy and cell survival. In addition, suppression of MAPK/ERK and TGF- β /Smad cascades contributes to reduced catabolic enzyme expression (MMP-13, ADAMTS-5) and fibrosis. Together, these pathways form a core regulatory network through which MSC-derived exosomal miRNAs exert anti-inflammatory, pro-regenerative, and cytoprotective effects, regardless of MSC tissue origin.

Prospects and Challenges in the Clinical Application of MSCs-Exosomal miRNAs in Arthritis

Based on previous preclinical studies, although cell-free exosomes have protective effects in OA/RA, their efficacy is only at early stages and several problems should be addressed. For example, the types and content of molecules in exosomes/EVs can be different depending on the tissue source, donor characteristics, and culture conditions of the parent MSCs. These points can alter exosomal miRNA profiles and consequently influence therapeutic potency and reproducibility. This point can impact MSC-exosomal function at recipient cells and cause alterations in physiological processes.

The efficiency of exosomal content delivery to chondrocytes is influenced by the exosome diameter (30–150 nm) because the average pore size of the articular cartilage ECM is about 6.0 nm⁸⁸ and only small cationic nanocarriers (<15 nm in diameter) can cross over this biological barrier.⁸⁹ During in vivo studies, the thickness of cartilage in small animal models is lower than humans⁹⁰ and cultured chondrocytes are investigated instead of full-thickness cartilage explants in most time during in vitro studies. These inconsistencies should be examined before the application of exosomes in the clinic.

The pleiotropic nature of miRNAs means that each can regulate multiple target genes; thus, off-target effects may occur if exosomal miRNAs inadvertently modulate pathways unrelated to joint repair, altering immune or metabolic functions in distant tissues. This concern is further heightened by evidence that repeated or high-dose administrations of MSC-derived exosomes could elicit unexpected immune activation or antibody formation, particularly if exosomes are derived from allogeneic sources. Moreover, key safety parameters such as dose-dependent toxicity and biodistribution remain poorly defined. Most preclinical studies use variable exosome concentrations and delivery routes without standardized pharmacokinetic evaluation. These issues make it difficult to predict or control these off-target and immunological risks.

The most important problem regarding the use of miRNAs is the delivery of appropriate amounts of selective miRNAs to save and repair cartilage tissue. The engineering of exosomes and their cargo, such as MSC pre-conditioning, exosome surface modification, and miRNA enrichment would be an effective therapy for RA and OA. Surface or genetic modification can enable targeted delivery of exosomes to inflamed synovial tissue or degenerative cartilage, thereby improving local uptake and minimizing off-target effects. Encapsulating MSC-derived exosomes within alginate hydrogels, collagen scaffolds, or chitosan-based matrices can allow sustained release and prolonged residence time at the target joint site. This point can address the rapid clearance typically observed after intra-articular injection.⁹¹ Future studies should therefore investigate optimal hydrogel formulations and release kinetics, as well as evaluate biocompatibility and functional outcomes in large-animal and clinical models. Comparative studies assessing exosomes derived from different MSC sources are warranted to determine which cell type yields the most functionally potent and clinically scalable exosomes. It is also important to investigate whether the complex regulatory network of miRNAs and their targets can initiate other diseases necessities further study.⁹²

Because of the local injection of exosomes mostly into the articular cavity, it is much safer than systematic administration and less toxic as cell-released products. Most in vivo MSC-exosome therapeutic studies were performed in small animals with less focus on their safety in arthritis treatment. Presently, insufficient evidence from clinical trials and preclinical studies delays the translation of MSC-exosomal therapies to clinical applications.

From a manufacturing and translational point, large-scale production of clinical-grade exosomes presents significant challenges (reviewed in Ref.⁹³). Scaling up MSC culture while maintaining cell viability, consistent phenotype, and exosome yield requires bioreactor-based culture systems and stringent Good Manufacturing Practice (GMP) compliance. Additionally, the lack of standardized protocols for exosome isolation and purification leads to variability in purity, particle concentration, and cargo integrity. Developing robust quality control assays to characterize exosomal size distribution, miRNA content, and bioactivity is essential for ensuring batch-to-batch consistency.

Exosome stability and storage also remain unresolved issues. Freeze–thaw cycles, lyophilization, and prolonged storage can alter exosomal membrane integrity and reduce biological activity.⁹³ Establishing validated methods for long-term preservation without functional loss will be critical for clinical manufacturing and distribution. Comparative profiling of MSC-derived exosomal miRNAs from healthy donors and arthritis patients, using sequencing or microarray approaches, may also identify disease-specific cargo signatures suitable for precision therapies. Addressing these challenges through standardized manufacturing protocols, rigorous dose optimization, and in vivo safety studies will be essential before clinical translation.

Conclusion

MSC-derived exosomal miRNAs represent a promising, low-risk, cell-free therapeutic strategy for OA and RA, combining the biological potency of stem cells with the safety and scalability required for clinical translation. In preclinical OA models, MSC-exosomal miR-92a-3p, miR-140-5p, miR-199a-3p, miR-95-5p, miR-320c and miR-100-5p are the most consistently reported therapeutic miRNAs. They protect cartilage and inflammation by targeting WNT5A/Wnt, ADAMTS/MMP, MAPK/NF- κ B, HDAC2/8 \rightarrow SOX9, ADAM-dependent Wnt and NOX4/mTOR (oxidative-stress/autophagy) pathogenic axes, respectively. In RA preclinical models, MSC-derived exosomes carrying miR-150-5p, miR-146a-5p, and miR-205-5p have demonstrated potent anti-inflammatory, anti-angiogenic, and joint-protective effects by downregulating key pathogenic mediators (MMP14, VEGF, MDM2) and reprogramming immune responses (eg, enhancing Treg, suppressing NF- κ B/MAPK).

Their intrinsic ability to cross biological barriers and their low immunogenicity make MSC-Exos an ideal platform for cell-free therapeutic development. Translationally, the identified miRNAs offer two major opportunities. First, as therapeutic targets, they can be enriched within exosomes through pre-conditioning of MSCs or encapsulated into advanced delivery systems to enhance joint repair and immune regulation. Second, as biomarkers, specific exosomal miRNA signatures may be used to monitor disease activity, predict treatment response, or stratify in future clinical trials. Future work should prioritize validating the most consistent miRNA candidates in human samples and developing standardized production and quantification methods to accelerate their path toward therapeutic and diagnostic application.

Abbreviations

ADAMTS9, ADAM metalloproteinase with thrombospondin type 1 motif 9; AVN, Avascular Necrosis or Osteonecrosis; BMSCs, Bone mesenchymal stem cells; CDH11, Cadherin-11; CXCL9, Chemokine ligand 9; DDX20, DEAD-Box Helicase 20; E2F2, E2F Transcription Factor 2; ELF3, E74 Like ETS Transcription Factor 3; FLS, Fibroblast-like synoviocyte; GREM1, Gremlin 1; IGF1R, Insulin-like growth factor1 receptor; hUC-MSCs, Human umbilical cord-derived mesenchymal stem cells; KLF, Krüppel-like factor; MAPK6, Mitogen-activated protein kinase 6; MDM2, Mouse double minute 2 homolog; MMP13, Matrix Metalloproteinase 13; mTOR, mechanistic Target of Rapamycin; MSC, Mesenchymal stem cell; NF- κ B, Nuclear Factor Kappa B; NOX4, NADPH Oxidase 4; OA, Osteoarthritis; PTGS2, prostaglandin-endoperoxide synthase 2; RA, Rheumatoid arthritis; ROS, Reactive oxygen species; SGK1, Serum/Glucocorticoid Regulated Kinase 1; hSMSCs, human synovial mesenchymal stem cells; TET1, Ten-eleven translocation methylcytosine dioxygenase 1; TGF- β , Transforming growth factor Beta; TNF- α , Tumor necrosis factor alpha; ADSCs: adipose-derived stromal cells, ATG4A: autophagy related 4A, GIT1: G-protein-coupled receptor kinase interacting protein-1.

Data Sharing Statement

Data will be made available upon a reasonable request.

Acknowledgments

The authors declare that they have not used AI-generated work in this manuscript.

Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

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Disclosure

The authors confirm that there is no conflict of interest related to the manuscript.

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