The basic characteristics of extracellular vesicles and their potential application in bone sarcomas

Shenglong Li1,2*

Abstract
Bone sarcomas are rare cancers accompanied by metastatic disease, mainly including osteosarcoma, Ewing sarcoma and chondrosarcoma. Extracellular vesicles (EVs) are membrane vesicles released by cells in the extracellular matrix, which carry important signal molecules, can stably and widely present in various body fluids, such as plasma, saliva and scalp fluid, spinal cord, breast milk, and urine liquid. EVs can transport almost all types of biologically active molecules (DNA, mRNA, microRNA (miRNA), proteins, metabolites, and even pharmacological compounds). In this review, we summarized the basic biological characteristics of EVs and focused on their application in bone sarcomas. EVs can be use as biomarker vehicles for diagnosis and prognosis in bone sarcomas. The role of EVs in bone sarcoma has been analyzed point-by-point. In the microenvironment of bone sarcoma, bone sarcoma cells, mesenchymal stem cells, immune cells, fibroblasts, osteoclasts, osteoblasts, and endothelial cells coexist and interact with each other. EVs play an important role in the communication between cells. Based on multiple functions in bone sarcoma, this review provides new ideas for the discovery of new therapeutic targets and new diagnostic analysis.

Highlights
New role of EVs in cell–cell communication in bone sarcoma.
New clinical practicality and future application prospects of EVs.
EVs can be use as biomarker vehicles for diagnosis and prognosis in bone sarcomas.
Bone sarcomas cells derived EVs may influence angiogenesis, osteoclastogenesis, immunomodulation, drug resistance, invasion, and migration processes.

Keywords: Extracellular vesicles, Bone sarcomas, Cancer diagnosis, Cancer therapy, Invasion and metastasis

Introduction
Bone sarcomas are malignant tumors that originates from mesenchymal tissue [1]. Osteosarcoma (OS) and chondrosarcoma are the most common malignant bone tumors, followed by Ewing sarcoma [1]. OS and Ewing sarcoma mainly occur in children and adolescents, while the incidence of chondrosarcoma increases with age [2, 3]. Osteosarcoma is the most common primary malignant bone tumor [4]. It mainly occurs in adolescents and children and reaches the second peak of incidence after old age [3]. Although neoadjuvant chemotherapy combined with surgical treatment can achieve a 5-year survival rate of 60–70%, for patients with relapsed and metastatic osteosarcoma, the original treatment method...
cannot produce effective therapeutic effects [5]. Chondrosarcoma is a malignant tumor derived from hyaline cartilage [6]. It can occur in any bone, but is common in the pelvis, humerus, femur, shoulder and ribs, and can occur at any age [6]. Patients with severe chondrosarcoma have a high mortality rate [7]. The treatment of chondrosarcoma includes aggressive surgical resection, systemic chemotherapy and targeted radiotherapy, but unfortunately, patients with chondrosarcoma often relapse and have a poor prognosis [7]. Ewing sarcoma mainly occurs in adolescents, and its malignant degree is high [8]. Recurrence and distant metastasis are the main causes of death [8]. Therefore, it is necessary to explore new treatment directions based on the development and metastasis mechanism of bone sarcomas.

Extracellular vesicles (EVs) are lipid bilayer nanovesicles secreted by cells, containing nucleic acids, proteins, lipids and other factors that maintain normal cell physiological functions and mediate cell-to-cell communication [9]. EVs are associated with multiple biological phenomena and are crucial for intracellular communication by transporting intracellular substances [10]. EVs are highly heterogeneous, and EVs secreted by different cells have different composition characteristics and functions. EVs have been regarded as wastes of cellular metabolism from the beginning to the current biological functions [11]. In addition to their important role in signal communication between cells, EVs are also widely involved in cell apoptosis, tumor development, angiogenesis, and immune response [12, 13]. Almost all cells secrete EVs under physiological and pathological conditions, and EVs can be found in blood, urine, saliva, and other body fluids [14]. EVs are widely observed in the tumor microenvironment [14]. They not only participate in the occurrence and development of tumors. Theoretically, tumor derived EVs are a dangerous "message in a bottle" for bone [15, 16]. EV plays a role in the regulation of bone remodeling activity and bone metastasis occurrence. They can modify the bone microenvironment, allowing the formation of osteolytic, osteosclerotic, and mixed metastasis [17]. However, the potential roles of EVs in the pathological exchange of bone cells between tumors and the bone microenvironment remain an emerging area [18]. The emerging evidence on EV functions in bone metastasis will facilitate the discovery of novel treatments [19].

In this review, we summarize the recent progress of the interaction between bone sarcoma and other cells in the tumor microenvironment through EVs, as well as the role of EVs as biomarker vehicles for diagnosis/prognosis and carriers for treatment in bone sarcomas. In particular, we discuss the role of these EVs in OS, Ewing sarcoma and chondrosarcoma, respectively. Compared with previous literature, we highlight the newly revealed role of EVs in cell–cell communication in bone sarcoma microenvironment, clinical practicality, and future application prospects.

### Biogenesis of EVs

EVs were first reported in 1946 as a platelet-derived procoagulant particles [20]. In 1983, a more detailed ultrastructural study found that during the differentiation of immature red blood cells, the fusion of multivesicular bodies (MVBs) with the cell membrane can also release similar vesicles and they are called exosomes [21]. EVs can be roughly divided into three categories: Microvesicles produced by budding and division from the plasma membrane. The intraluminal vesicles released when MVBs fuse with the plasma membrane, namely exosome [22]. Apoptotic bodies released in the form of cell vesicles during cell apoptosis [23]. The size of EVs is usually used as the classification criteria: small vesicles < 150 nm are classified as exosomes, and those with 100~1000 nm or more are classified as micro-vesicles [24] (Fig. 1). There is still a lack of consensus on the nomenclature of extracellular vesicles. The International Society of Extracellular Vesicles (ISEV) encourages the use of "extracellular vesicles" as a general term and key word for all secreted vesicles [25]. The formation and secretion of EVs depends on the participation of endosomal sorting complex required for transport (ESCRT) [26]. As a consequence of their origin, exosomes from different cell types contain endosome-associated proteins (e.g., Rab GTPase, SNAREs, Annexins, and flotillin), some of which are involved in MVB biogenesis (e.g., Alix and Tsg101) [27]. The way in which virus-like microvesicles sprout on the surface of cell membrane is usually used as the classification criteria: small vesicles < 150 nm are classified as exosomes, and those with 100~1000 nm or more are classified as micro-vesicles [24] (Fig. 1). There is still a lack of consensus on the nomenclature of extracellular vesicles. The International Society of Extracellular Vesicles (ISEV) encourages the use of "extracellular vesicles" as a general term and key word for all secreted vesicles [25]. The formation and secretion of EVs depends on the participation of endosomal sorting complex required for transport (ESCRT) [26]. As a consequence of their origin, exosomes from different cell types contain endosome-associated proteins (e.g., Rab GTPase, SNAREs, Annexins, and flotillin), some of which are involved in MVB biogenesis (e.g., Alix and Tsg101) [27]. The way in which virus-like microvesicles sprout on the surface of cell membrane is usually used as the classification criteria: small vesicles < 150 nm are classified as exosomes, and those with 100~1000 nm or more are classified as micro-vesicles [24] (Fig. 1). There is still a lack of consensus on the nomenclature of extracellular vesicles. The International Society of Extracellular Vesicles (ISEV) encourages the use of "extracellular vesicles" as a general term and key word for all secreted vesicles [25].

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### Protein: Most of the proteins contained in EVs are shared by different types of EVs, such as tetrameric proteins (CD9, CD63, CD81 and CD82), 14–3–3 proteins, Major histocompatibility complex (MHC) molecules and specific stress proteins (heat shock proteins) and other cytoplasmic proteins; Endosome sorting
complex 3 (ESCRT-3) binding protein required for transport. In general, EVs are very rich in cytoskeleton proteins, cytoplasmic proteins, heat shock proteins, cell membrane proteins, and proteins involved in vesicle transport, while there are fewer organelle proteins in the cell [30].

Lipids: EVs are different from their secreting cells in terms of lipid composition, and there may be a mechanism that can classify these specific lipid types into the vesicle [31].

Nucleic acid: EVs contain complete mRNA, mRNA fragments, long non-coding RNA (lncRNA), miRNA, ribosomal RNA (rRNA), different EVs may have different types and levels of these Nucleic acids [13]. EVs are loaded with content molecules of different types and contents to reflect the different states of parent cells. These substances also affect the properties and functions of EVs. For example, the content mRNA loaded by them is transferred horizontally through EVs and enters the recipient cells. It is translated into protein to change the biological state and function of the recipient cell [32], while miRNA can be stored in EVs in the blood circulation to avoid the degradation of RNAse, and then combine with immune cells to play an immunomodulatory effect. The system is an indispensable and important part in the occurrence and development of tumors [33].

The detection methods of EVs
After the operation of separating and purifying extracellular vesicles, the morphology and purity of the extracellular vesicles need to be detected before sequencing or protein profiling of their contents (Table 1). This is a necessary condition to ensure the reliability of the later analysis data.

Electron microscope inspection
Electron microscopy is the gold standard for morphological detection of extracellular vesicles [34, 35]. The resolution of the transmission electron microscope is 1 ~ 3 nm, and the resolution of the scanning electron microscope is 5 nm. When observing an electron microscope sample, the diameter of extracellular vesicles can be measured. The immunogold labeling method can be used to label proteins on the surface of extracellular vesicles. Cryo-electron microscopy can observe the double-layer membrane structure of extracellular vesicles in order to distinguish between extracellular vesicles and other non-vesicular structures. Electron microscopy can reveal the structure of purified single extracellular vesicles or apoptotic bodies in tissues [36].

Flow cytometry
Flow cytometry is an important method to analyze extracellular vesicles [34, 35]. Flow cytometric analysis sorts
| Classification                  | Detection technology                  | Advantages                                                                                   | Disadvantages                                                                 |
|--------------------------------|---------------------------------------|------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------|
| Separation and enrichment      | Ultracentrifugation                   | Simple operation, can be used for large quantities of samples                                | The equipment required is expensive and time-consuming                          |
|                                | Density gradient centrifugation       | The separation purity is improved, and the EV activity can be better maintained               | Time-consuming, the osmotic pressure must be controlled when preparing inert gradient media |
|                                | Size exclusion chromatography         | The separation purity is further improved, which saves time and can better maintain EV activity| The number of times the column is used and the amount of sample loaded are limited, and lipoproteins in some samples may be co-separated with EV |
|                                | Ultrafiltration                       | Simple operation, time-saving, can better maintain EV activity                                | Cannot filter out impurities smaller than the pore size of the filter membrane   |
|                                | Polyethylene glycol precipitation method | Reagents are easy to get, and the operation is simple                                          | Time-consuming and susceptible to interference from other hydrophobic proteins, affecting the subsequent protein functional analysis of EV |
|                                | Immunomagnetic bead sorting           | Good specificity and high purity                                                              | Higher cost, limited by antibody preparation technology; epitope can be activated or blocked, affecting subsequent functional analysis |
| Identification                | Electron microscope inspection        | Can be used for EV morphological characterization                                             | High equipment requirements, complicated sample preparation, dehydration, fixation, and dyeing may affect EV activity and high cost |
|                                | Nanoparticle tracking analysis technology | Real-time EV concentration and particle size distribution information can be obtained          | Unable to distinguish the EV phenotype from the source, and the EV cannot be distinguished from particles of similar size; not suitable for heterogeneous samples, the light intensity signal of large particles can easily mask the signal of small particles |
|                                | western blot and Elisa                | It can perform qualitative, semi-quantitative or quantitative analysis of the target protein with strong specificity | Only known proteins can be detected, limited by antibody preparation technology, the operation is cumbersome and time-consuming |
of particles according to the size of extracellular vesicles or the fluorescent signals carried. The size of extracellular vesicles is mainly analyzed by its scattered light, and the fluorescence carried by extracellular vesicles is mainly analyzed by the emission light of extracellular vesicles under laser excitation. The intensity of light scattered by extracellular vesicles is weak, sometimes lower than background noise. Extracellular vesicles can be labeled with fluorescent antibodies to be detected in flow cytometry. However, due to the small size of single extracellular vesicles, the abundance of labeled proteins on the surface is low, so the fluorescence intensity is lower than that of most streams. The resolution of the cytometer. Therefore, flow cytometry needs to distinguish the signal of extracellular vesicles under a lot of background noise.

Nanoparticle tracking analysis technology
One of the most common methods for identifying EVs based on particle size is Nanoparticle Tracking Analysis Technology (NTA) [35]. This technology is to install a high-definition camera on an optical microscope, using the properties of light scattering and Brownian motion, through the Stokes-Stein equation (the movement speed of nanoparticles in their suspension per unit time and their own particle size). There is a quantitative relationship between the viscosity of the solution and the temperature, the specific exosomes and microvesicles in the diameter range of 50 ~ 1000 nm are directly imaged and observed one by one, and the high-resolution particle size distribution data and concentration are obtained.

Antibody-based identification method
Given that EV is produced in the cell membrane pathway, antibody targeting markers related to this pathway can be identified [35, 37]. These include the four transmembrane protein superfamilies (CD9, CD63 and CD81), AIP1/Alix, TSG101 and CD326/EPCAM [38]. The method of identification can be western blotting.

Fluorescence and Confocal Microscopy
EV can be labeled with lipophilic membrane-bound dyes (such as PKH67, DiD, etc.), or the sulphydryl group on its surface can be used to label EVs [39, 40]. This technology cannot really visualize each EV, but it can be used to study whether the labeled EV can be taken up by cells.

The role of EVs in osteosarcoma
As a medium, EVs play a vital role in the communication between tumor cells and other cells in the tumor microenvironment (Fig. 2). Bone sarcomas cells can interact with surrounding cells through input/output of EVs. EV-mediated crosstalk occurs through the trafficking of vesicle-associated components to endothelial cells, osteoclasts, T cells, CTCs, CAFs, MSCs, and bone sarcomas cells. CAFs may transfer the ability of migration/invasion through EVs to bone sarcomas cells. Neighboring stem cells may transfer factors contributive to growth and metastasis. Bone sarcomas cells derived EVs may influence angiogenesis, osteoclastogenesis, immunomodulation, drug resistance, invasion, and migration processes. The roles of EVs in different types of bone sarcoma are as follow.

The role of EVs in osteosarcoma
OS is the most common primary bone tumor, which occurs in 0.3 cases per 100,000 people [41]. Although diagnosis and treatment have improved in the past few decades, the survival rate for a considerable number of patients is still very low [3]. Therefore, one of the focuses of OS research is to better understand the metastatic process and how different factors regulate MTCT to promote metastatic spread. This will help determine treatment strategies for metastatic and refractory diseases, thereby improving survival.

Since EVs play a role in the tumorigenesis of many cancers and the role of these small extracellular vesicles in metastasis, it is not surprising that researchers in the OS field are studying the role of EVs in OS (Table 2).

Studies have shown that EVs derived from other cells can affect the function of osteosarcoma cells. Exosomes derived from bone marrow mesenchymal stem cells (BMSCs) could transport miRNA 206 (miR-206) to osteosarcoma cells. Mechanically, exosomal-miR-206 may inhibit the proliferation, migration and invasion of osteosarcoma cells by targeting TRA2B, and it may induce OS cell apoptosis [42]. Zhao et al. proved that BMSCs-derived exosomes encapsulated long non-coding PVT1 RNA and transported it to osteosarcoma cells, and the transported PVT1 promoted tumor growth by inhibiting ubiquitination and promoting ERG expression in osteosarcoma cells [43]. In addition, Qin et al. had shown that BMSC-derived exosomes miR-208a can improve the progression of osteosarcoma by targeting PDCD4 [44]. Wang et al. found that adipose-derived mesenchymal stem cells (AD-MSCs) exosomes can promote the progression of osteosarcoma by targeting PDCD4. Exosomes derived from bone marrow mesenchymal stem cells (BMSCs) could transport miRNA 206 (miR-206) to osteosarcoma cells. Mechanically, exosomal-miR-206 may inhibit the proliferation, migration and invasion of osteosarcoma cells by targeting TRA2B, and it may induce OS cell apoptosis [42]. Zhao et al. proved that BMSCs-derived exosomes encapsulated long non-coding PVT1 RNA and transported it to osteosarcoma cells, and the transported PVT1 promoted tumor growth by inhibiting ubiquitination and promoting ERG expression in osteosarcoma cells [43]. In addition, Qin et al. had shown that BMSC-derived exosomes miR-208a can improve the progression of osteosarcoma by targeting PDCD4 [44]. Wang et al. found that adipose-derived mesenchymal stem cells (AD-MSCs) exosomes can promote the progression of osteosarcoma by increasing the expression of COLGALT2 in osteosarcoma cells [45]. Human exosomes derived from BMSCs might promote the growth and metastasis of OS by promoting oncogenic autophagy [46]. Ge et al. demonstrated that BMSCs-derived exosomes LCP1 could promote bone proliferation and metastasis through JAK2/STAT3 pathway [47]. The targeting of miR-135a-5p/LCP1 axis might have potential in the treatment of OS [47]. The macrophage-derived exosomes Inc-LIFR-AS1 could promote the proliferation, invasion, and apoptosis of osteosarcoma cells.
through the miR-29a/NFIA regulatory axis [48]. Cancer-associated fibroblasts (CAFs) could spread exosomal miR-1228 by targeting SCAI, thereby promoting the invasion and migration of OS [49]. AD-MSCs could target BCL-6 to obtain miR-101-rich exosomes in OS cells, thereby inhibiting tumor growth and metastasis [50].
In addition, bone sarcomas can also secrete EVs to regulate their own tumor growth and metastasis. The results of Gong et al. study showed that metastatic OS cells could transfer exosomal miR-675 to non-metastatic cells and promoted cell migration and invasion by targeting CALN1 [51]. The inactivation of Hic-5 could inactivate Wnt/β-catenin signal through exosomal pathway, thereby inhibiting proliferation and inducing apoptosis of osteosarcoma cells [52]. The exosomal miR-1307 from OS cells promoted the proliferation, migration and invasion of OS cells by targeting AGAP1, and the miR-1307-AGAP1 axis might play an important role in the future treatment of OS [53]. In addition, the exosomal linc00852 associated with AXL up-regulated the proliferation, migration and invasion of OS cells, which was considered a novel tumor biomarker and a special therapeutic target for OS [54]. Towards endothelial cells, OS-derived exosomes promote osteoclasts differentiation and bone resorption activity, and these exosomes potentiated tube formation of endothelial cells and increased angiogenic markers expression. The molecular mechanisms underlying this process may include miR-148a and miR-21-5p [55, 56].

Studies have also shown that OS derived EVs might also participate in bone development by affecting the function of surrounding cells. Compared with normal osteoblasts, exosomes from osteosarcoma contain immunomodulatory substances, which could reduce the proliferation rate of T cells and promote the regulatory phenotype T [57]. Osteoblast exosomes could also reduce T cell activity, but to a lesser degree than canine osteosarcoma (OSA) exosomes and do not promote the T regulatory phenotype [57]. OS-derived exosomes might induce M2 polarization of macrophages and promote the invasion and metastasis of tumors by Tim-3 [58]. Bone sarcomas derived exosomal miR-501-3p promoted osteoclast production and aggravate bone loss through the PTEN/PI3K/Akt signaling pathway [59]. The Rab22a-NeoF1 fusion protein is secreted into exosomes through its KFERQ-like motif binding to HSP90. Macrophages and cancer cells negative for the fusion gene absorb the protein, while the exosomal fusion protein Rab22a-NeoF1 could promote its receptor-negative cancer cells metastasize in mouse lungs through the activation of RhoA activation by the binding partner PYK2 of their donor cells [60]. OS-derived EVs could recapitulate the infiltration

**Table 2** Summary of EVs studies in OS

| Cargos          | Parent cell | Target cell | Biological function                                                                 | Reference |
|-----------------|-------------|-------------|-------------------------------------------------------------------------------------|-----------|
| miR-206         | BMSCs       | OS cells    | Cell proliferation, migration, invasion and apoptosis                                 | [42]      |
| Lnc-PVT1        | BMSCs       | OS cells    | Tumor growth                                                                         | [43]      |
| miR-208a/PCD4   | BMSCs       | OS cells    | Promote OS progression                                                               | [44]      |
| COLGALT2        | AD-MSCs     | OS cells    | Promote OS progression                                                               | [45]      |
| miR-135a-5p/LCP1| BMSCs       | OS cells    | Promote OS proliferation and metastasis                                             | [46]      |
| miR-LIFR-AS1/miR-29a/NFIA | macrophage | OS cells    | Cell proliferation, migration, invasion and apoptosis                                 | [47]      |
| miR-1228/SCAI   | CAFs        | OS cells    | Cell proliferation, migration, invasion and apoptosis                                 | [48]      |
| miR-151-3p/CHL1/Integrin 1β | CAFs | OS cells    | Cell proliferation, migration, invasion and apoptosis                                 | [49]      |
| miR-101/BCL-6   | AD-MSCs     | OS cells    | Tumor growth                                                                         | [50]      |
| miR-675/CAN1    | metastatic OS cells | Non-metastatic OS cells | Tumor growth and metastasis                                                        | [51]      |
| Hic-5/Wnt/β-catenin | OS cells | OS cells | Cell proliferation and apoptosis                                                      | [52]      |
| miR-1307/AGAP1  | OS cells    | OS cells    | Cell proliferation, migration, and invasion                                          | [53]      |
| miR-S01-3p      | OS cells    | Osteoclast  | Promote osteoclast production and aggravate bone loss                                | [54]      |
| LINE-1          | OS cells    | CAFs        | Cell differentiation                                                                 | [55]      |
| TGFB2           | Metastatic OS cells | MSCs | Epigenetic transformation                                                           | [56]      |
| COL6A1          | OS cells    | CAFs        | Activate CAF to promote OS transfer                                                 | [57]      |
of myeloid cells into the lungs of naive mice, but it is not enough to promote OS metastasis [61]. Mazumdar et al. also proved that EVs derived from OS cells could cause cancer-related fibroblast/fibroblast differentiation [62]. The OS cell line was able to produce EVs fused with recipient cells, and under the conditions of starvation, high-level activation of survival pathways, migration, adhesion, and 3D enhancement of ball formation, enhance its ability to grow in anchors, thereby enhancing proliferation and survivability [63]. OS-exosome-mediated LINE-1 methylation was insufficient in MSCs, while the opposite effect was observed in osteoblasts, indicating that MSCs are sensitive to epigenetic transformation but not to osteoblasts [64]. The exosomes of metastatic osteosarcoma cells could regulate the cell signaling of tumor-associated macrophages, thereby promoting the M2 phenotype by producing TGFβ2 and creating an immunosuppressive microenvironment that promotes tumors [65]. Zhang et al. proved that COL6A1 can be packaged in OS derived exosomes and activate CAFs to promote OS transfer [66].

The role of EVs in Ewing sarcoma
Ewing sarcoma (EWS) is a malignant tumor commonly seen in children and adolescents [67]. The only prognostic factor for patients with recurrence is the recovery time. Those who relapse 2 years after the initial diagnosis have a relatively good prognosis [68]. The 5-year survival rate of patients with local recurrence is 13%-30%, but the prognosis of patients with systemic or other tumor recurrence is better [68]. Recent studies have shown that EVs also play an important role in the development of Ewing sarcoma.

Feo et al. had shown that the elimination of CD99 in EWS tumor cells leads to the production and release of exosomes. These exosomes could transfer their anti-tumor effects to neighboring tumor cells. This indicated that these exosomes are in the reversal of malignant tumors rather than initiation in the soil. An atypical new role was played in the process transfer seeding [69]. Ventura et al. confirmed that the delivery of exosomes through CD99-silenced cells was sufficient to inhibit Notch-NF-kB signaling via miR-34a to induce neural differentiation of recipient EWS cells [70]. Miller et al. proved that EWS-derived exosomes might be used as biomarkers to minimize the diagnosis of residual diseases in peripheral blood, and prompt people to further study their potential biological effects in modifying the microenvironment related to EWS [71]. Hypoxic exosomes promoted stems in EWS cells by providing enriched miR-210 that could down-regulate the apoptotic pathway, leading to cell survival and increasing sphere formation [72].

The role of EVs in chondrosarcoma
Chondrosarcoma is a malignant tumor that originates from cartilage or cartilage-forming connective tissue [3]. The incidence of malignant bone tumors ranks second, second only to osteosarcoma. The clinical manifestations of most lesions (especially secondary) are slow development, long duration, mild symptoms, and good prognosis; a few lesions (especially primary) progress fast, short duration, severe symptoms, and poor prognosis [3]. Cheng et al. found that chondrosarcoma cell-derived exosomes carry IncRNA RAMP2-AS1 and regulate the angiogenic ability of HUVECs via acting as a ceRNA of miR-2355-5p to regulate VEGFR2 expression [73].

EVs as biomarker vehicles for diagnosis and prognosis in bone sarcomas
EVs are widely present in almost all body fluids, containing nucleic acids, proteins, lipids, metabolites, etc. Under different cell sources and different physiological or pathological conditions, the composition and content of EVs content will change significantly, and the level of specific content will change. The detection can reflect the physiological and pathological state of cells, and has the potential of liquid biopsy markers. At present, a variety of research strategies have been used in the screening of EVs markers, each with its advantages and limitations, and seeking the best clinical research strategy is still the key to screening for markers with high application value.

Detection of PD-L1 and exosomal N-cadherin in the serum of OS patients could predict the progression of lung metastasis in OS patients [74]. Cambier et al. verified that a consistent excess of DNA sequences of repetitive elements associated with EVs indicates their potential use as biomarkers for OS [75]. Li et al. found that SENP1 derived from plasma exosomes could be used as a new independent prognostic indicator in the clinical application of OS [76]. Zhang et al. introduced the latest progress of EWS and the opportunities and challenges brought by the development of circulating exosomes as a diagnosis and monitoring of children and young adults in the EWS family (ESFT) source of development of biomarkers for treatment response in adult patients [77]. Samuel and colleagues had also shown that circulating EVs could be used as a source of minimally invasive and potential prognostic diagnostic biomarkers in pediatric patients with tumors [78]. In addition, compared with the control, CASC15 upregulation was observed in OS plasma exosomes, and the same expression was observed in OS tissues and cell lines [79]. Besides, 30 gene fusions related to cancer patients have been identified as events in EVs RNA and are more common in metastatic EVs [80]. Analysis strategies for serum exosomal miRNAs and mRNAs have been developed for OS patients with
different chemotherapeutic responses [81]. Compared with OS patients with good chemotherapy response, 12 miRNAs in OS patients with poor chemotherapy response were up-regulated, while 18 miRNAs were significantly down-regulated [81].

The application of EVs in bone sarcoma treatment

The use of EVs to treat human diseases has become a core issue in clinical medicine because of their ability to deliver biologically active substances to target cells. Therefore, EVs are regarded as natural nanocarriers with high therapeutic potential [42, 82–86].

Pan et al. showed that exosomes from cisplatin-resistant cells (CDDP) reduced the sensitivity of MG63 and U2OS cells to CDDP, inhibited cell apoptosis, and increased the levels of multidrug resistance-related protein 1 and P-glycoprotein expression [87]. In addition, exosomal hsa_circ_103801 could enhance the promoter function of exosomes and promote the chemoresistance of MG63 and U2OS cells to CDDP [87]. Wei et al. found that the prepared doxorubicin-loaded exosomes could be used as an excellent chemotherapeutic drug for the treatment of osteosarcoma in vitro [88]. Considering the tumor localization function of BM-MSCs, doxorubicin-loaded exosomes might be a novel candidate for targeted therapy of OS in future studies. In addition, multidrug-resistant OS cells could expand their ability to resist the effects of adriamycin on sensitive cells by transferring exosomes carrying MDR-1 mRNA and its P-glycoprotein product [89].

Remaining concerns and future perspectives

EVs are widely found in organisms, and their biological functions are increasingly recognized [90, 91]. As a natural communication medium between cells, EVs are expected to be used to treat a variety of clinical diseases based on this feature [11]. In addition, due to their high bioavailability and low immunogenicity, they can be the best candidates for drugs and therapeutic molecular carriers [12, 92]. For example, heterologous exosomes released by mesenchymal stem cells are considered a reliable and safe source of therapeutic exosomes [93, 94]. Clinically, the use of autologous methods is not ruled out, because in this case, the possibility of exosomes containing potentially dangerous molecules is very small. However, the exosomes of the patient’s plasma are dangerous. Because the molecules delivered by the plasma exosomes are considered to be some metabolic wastes of diseased tissues, they are likely to deliver some drugs with high toxic potential. These studies support the use of exosomes secreted by primary monocytes in peripheral blood as drug carriers. Of course, it is not to say that they are completely safe, but they are safer than plasma exosomes, which requires clinical research to determine the true therapeutic potential of EVs. However, the most likely problem to be solved is the processing of EVs content. To this end, the best results can be achieved by establishing “EVs factories” similar to cell therapy cell factories. Another issue that needs to be addressed is the mechanism by which EVs play a therapeutic role. The mechanism of the interaction between EVs and target cells may be membrane-membrane fusion or the delivery of vesicle contents in target cells. Of course, they themselves may trigger effects. For example, EVs secreted by different types of cells may preferentially target certain cell types depending on the composition of the membrane, thereby having different effects on our body. Nevertheless, the mechanism of how EVs play a therapeutic role and affect target cells remains to be elucidated. Whether it is direct modification of EVs or selection of different cell sources for EVs, the safety issues still exist. Therefore, the research in the next few years may focus on the research on the impact of EVs on the body. There is no doubt that these goals can only be achieved after careful and in-depth research on the content and characteristics of EVs that have not been used in clinical practice.

In the microenvironment of bone sarcoma, bone sarcoma cells, mesenchymal stem cells, immune cells, fibroblasts, osteoclasts, osteoblasts, and endothelial cells coexist and interact with each other [15, 95, 96]. EVs play an important role in the communication between cells. On the one hand, osteosarcoma cells secrete EVs to reach the recipient cells to promote tumor support properties. On the other hand, EVs derived from tumor microenvironment cells can help tumor growth and migration. EVs with specific cell membrane components and specific wraps of the source cells have great potential as diagnostic and prognostic markers. EVs have the above-mentioned multiple functions in bone sarcoma, providing new ideas for the discovery of new therapeutic targets and new diagnostic analysis.

In the future, technological advances in the purification and characterization of EVs are expected to better help the detection of EVs and the study of their biological characteristics, and its clinical application prospects in bone sarcoma will be broader. At least, future research can focus on the following aspects. First, the identification of the expression profile of EV-specific inclusions is helpful for machine learning to identify the occurrence and types of OS. Second, the mechanism of transmission of drug resistance or metastatic properties by EVs can be further explored from multiple dimensions. Third, which type of EV is most easily ingested by bone sarcoma is a question to be addressed for the development of targeted drug carriers.
Abbreviations
EVs: Extracellular vesicles; miRNAs: MicroRNAs; OS: Osteosarcoma; EWS: Ewing sarcoma; MVBs: Multivesicular bodies; ISEV: International Society of Extracellular Vesicles; ESCRT: Endosomal sorting complex required for transport; MHC: Major histocompatibility complex; IncRNA: Long non-coding RNA; rRNA: Ribosomal RNA; PEG: Polyethylene glycol; NTAs: Nanoparticle Tracking Analysis; BMSCs: Bone marrow mesenchymal stem cells; ERG: ETS transcription factor ERG; PDCD4: Programmed cell death 4; AD-MSCs: Adipose-derived mesenchymal stem cells; COLGALT2: Collagen beta(1-O)galactosyltransferase 2; LCP1: Lymphocyte cytosolic protein 1; NFIA: Nuclear factor I A; CAFs: Cancer-associated fibroblasts; CHL1: Cell adhesion molecule L1 like; CALN1: Calneuron 2; AGAP1: ArfGAP with GTPase domain, ankyrin repeat and PH domain 1; COL6A1: Collagen type VI alpha 1 chain.

Acknowledgements
This work was supported by the Liaoning Cancer Hospital & Institute (Shenyang) and China Medical University (Shenyang).

Authors’ contributions
Original draft preparation, allocation, revision, supplement and edition: SL. All authors have read and agreed to the published version of the manuscript.

Funding
This work was supported by Natural Science Foundation of Liaoning Province (2020-MS-058) and Shenyang young and middle-aged scientific and technological innovation talent support plan (RC190456).

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

Declarations
Ethics approval and consent to participate
None.

Consent for publication
None.

Competing interests
The authors declare no conflict of interest.

Author details
1 Department of Bone and Soft Tissue Tumor Surgery, Cancer Hospital of China Medical University, Liaoning Cancer Hospital & Institute, Shenyang 110042, Liaoning Province, China. 2 Department of Tissue Engineering, Center of 3D Printing & Organ Manufacturing, School of Intelligent Medicine, China Medical University (CMU), No. 77 Puhe Road, Shenyang North New Area, Shenyang 110122, China.

Received: 15 May 2021 Accepted: 7 September 2021
Published online: 17 September 2021

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