

Bone Marrow Mesenchymal Stem Cell-derived Exosomes Exert Therapeutic Effects by Attenuating Fibrosis and Facilitating Hepatocyte Proliferation via the Anti-pyroptosis Pathway in a Rat Cirrhosis Model

Yichi Zhang

Second Affiliated Hospital of Harbin Medical University

Xinsheng Nie

xinjiang shengchan jianshe bingtuan: Xinjiang Production and Construction Corps

Lijuan Wang

xinjiang shengchan jianshe bingtuan: Xinjiang Production and Construction Corps

Zhuzhi Wan

xinjiang shengchan jianshe bingtuan: Xinjiang Production and Construction Corps

Hao Jin

xinjiang shengchan jianshe bingtuan: Xinjiang Production and Construction Corps

Ronghui Pu

xinjiang shengchan jianshe bingtuan: Xinjiang Production and Construction Corps

Meihui Liang

xinjiang shengchan jianshe bingtuan: Xinjiang Production and Construction Corps

Yuan Chang

Second Affiliated Hospital of Harbin Medical University

Yang Gao

Second Affiliated Hospital of Harbin Medical University

Hailong Zhang

xinjiang shengchan jianshe bingtuan: Xinjiang Production and Construction Corps

Shizhu Jin (✉ drshizhujin@hrbmu.edu.cn)

Second Affiliated Hospital of Harbin Medical University <https://orcid.org/0000-0003-3613-0926>

Research

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1 **Bone marrow mesenchymal stem cell-derived exosomes exert**
2 **therapeutic effects by attenuating fibrosis and facilitating hepatocyte**
3 **proliferation via the anti-pyroptosis pathway in a rat cirrhosis model**

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5 Zhang YC¹, Nie XS², Wang LJ², Wan ZZ², Jin H², Pu RH², Liang MH², Chang Y¹,
6 Gao Y¹, Zhang HL², Jin SZ^{1,2*}.

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8 ¹ Department of Gastroenterology and Hepatology, The Second Affiliated Hospital,
9 Harbin Medical University, Harbin, Heilongjiang Province, China.

10 ² The Xinjiang Production and Construction Corps Tenth Division Beitun Hospital,
11 Beitun, Xinjiang Province, China.

12

13 * Corresponding author. Correspondence and requests for materials should be
14 addressed to Shizhu Jin. (email: drshizhujin@hrbmu.edu.cn)

15

16 **Abstracts**

17 **Background:** Liver cirrhosis, which lacks a specific therapy, induces a substantial
18 economic burden and affects quality of life. Exosomes are nanosized vesicles that are
19 secreted by cells transfer bioactive molecules to regulate metastasis and have attracted
20 immense interest as a therapeutic option for liver cirrhosis. Therefore, in this study,
21 we explored the efficacy and dose of exosomes derived from rat bone marrow
22 mesenchymal stem cells (Exos-rBMMSC) in a hepatic cirrhosis rat model.

23 **Results:** Four weeks after exosome therapy, NLRP3, GSDMD, cleaved caspase-1,
24 IL-1 β and IL-18 expression levels in the 200 μ g and 400 μ g Exos-rBMMSC groups
25 were significantly decreased compared to those in the liver cirrhosis group ($P < 0.05$).
26 Similar results were observed in the hepatic function assay; serum levels of aspartate
27 aminotransferase and alanine aminotransferase were decreased, and albumin was
28 increased. The histopathology results showed that 200 μ g and 400 μ g of
29 Exos-rBMMSC significantly relieved collagen deposition, while 100 μ g of
30 Exos-rBMMSC induced no marked improvements compared to those in the liver
31 cirrhosis group. The localization of PKH67-labelled Exos-rBMMSC was verified
32 microscopically, and these particles coexpressed PCNA, NLRP3, GSDMD and
33 Caspase-1.

34 **Conclusions:** Our findings suggest that Exos-rBMMSC accelerate hepatocyte
35 proliferation and relieve liver fibrosis, which may be a promising therapeutic strategy
36 for liver cirrhosis.

37 **Key words:** Exosomes; BMMSCs; Liver cirrhosis; Pyroptosis

38

39 **Introduction**

40 Liver cirrhosis is the terminal phase of all types of chronic hepatic diseases.
41 Currently, without an effective therapy for liver cirrhosis, patients are more likely to
42 experience hospital readmissions, hepatocellular carcinoma and death, which results
43 in a substantial economic burden. The annual death rate of liver cirrhosis worldwide is
44 as high as 1.32 million [1]. Therefore, developing an effective therapy is an urgent
45 problem to be solved.

46 In recent years, accumulating evidence from preclinical experiments and clinical
47 trials has demonstrated that rat bone marrow mesenchymal stem cell (rBMMSC)
48 transplantation can alleviate collagen accumulation and improve liver function [2, 3].
49 However, the therapeutic effect is limited by low differentiation efficiency, and most
50 stem cells are obstructed in the pulmonary vasculature [4]. Moreover, the clinical
51 application of stem cell transplantation is hampered by complications such as
52 tumorigenicity and immunologic rejection [5]. Hence, exosomes (Exos), which
53 possess similar biological topology as stem cells, are important paracrine mediators
54 and have become a promising and advanced cell-free remedy. Exos are vesicles that
55 are 30~150 nm in diameter, are secreted by all kinds of cells, and transfer functional
56 mircoRNAs (miRNAs), mRNAs, proteins and cytokines, and these vesicles have
57 inspired substantial research enthusiasm for mediating intercellular metastasis [6; 7].
58 However, few studies have focused on the applicable doses of Exos, which is a gap
59 that hampers practical clinical applications.

60 In this study, we aimed to explore the optimal dose of Exos derived from
61 rBMMSCs (Exos-rBMMSC) for treating liver cirrhosis. The therapeutic effects of
62 100 µg, 200 µg and 400 µg of Exos-rBMMSC were evaluated by examining
63 pyroptosis inhibition, pathological histology, liver function and molecular biology in
64 vivo and in vitro. In addition, the distribution of Exos was examined by near-infrared
65 fluorescence (NIRF) imaging. Overall, our results suggested that Exos-BMMSC
66 alleviated liver cirrhosis and enhanced hepatic function.

67

68 **Results**

69 Liver cirrhosis model establishment.

70 Rat liver cirrhosis models were verified by analysing specimens, and the results
71 showed that the surface of the model liver was grainy and pale, and the organ was
72 smaller in size than the control liver (Fig. 1A). Haematoxylin and eosin (HE)-stained
73 cirrhotic liver sections showed cellular necrosis, inflammatory cell infiltration and
74 pseudolobuli formation compared to those in the control group (Fig. 1B & 1D).
75 Masson staining of sections showed that compared to those in the control group,
76 hepatic collagen production and collagen deposition were obviously present (Fig. 1C
77 & 1E).

78 rBMMSCs alleviated hepatocyte pyroptosis.

79 We confirmed that pyroptosis-related proteins were highly expressed in the liver
80 cirrhosis model (Fig. 1F). In addition, 4 weeks of rBMMSC transplantation via caudal
81 vein injection significantly alleviated the degree of liver fibrosis (Fig. 1G & 1H) and
82 markedly restrained the expression of pyroptosis-related proteins (NLRP3, GSDMD
83 and cleaved caspase-1, Fig. 1I-1M, * $P < 0.05$). IL-1 β and IL-18, which are
84 pyroptosis-related inflammatory factors, were decreased after 4 weeks of rBMMSC
85 transplantation (Fig. 1N & 1O, ** $P < 0.01$).

86 Paracrine effect of rBMMSCs on inhibiting hepatocyte pyroptosis.

87 After hepatocytes and rBMMSCs were cocultured for 24 h, PKH67 dye was
88 observed in hepatocytes, which indicated that rBMMSCs affected hepatocytes in a
89 paracrine manner (Fig. 2A-2D). GW4869 (dissolved in DMSO, 10 μ M) is an inhibitor
90 of membrane neutral sphingomyelin, which participates in Exo generation and
91 release[8]. To assess the effect of Exos-rBMMSC on hepatocytes, the following
92 groups were established: control group (untreated hepatocytes), CCl₄ group
93 (hepatocytes were treated with 10 mM CCl₄ (dissolved in DMSO)), CCl₄+DMSO
94 group (hepatocytes were treated with CCl₄ (10 mM) and 1% DMSO), CCl₄+GW4869
95 group (hepatocytes were treated with CCl₄ (10 mM) and GW4869 (10 μ M)),
96 CCl₄+CCS group (hepatocytes were treated with CCl₄ (10 mM) and rBMMSC cell
97 culture supernatant (CCS)), and CCl₄+CCS+GW4869 group (hepatocytes were

98 treated with CCl₄ (10 mM), GW4869 (10 μM) and rBMMSC CCS).
99 Pyroptosis-related protein (NLRP3, GSDMD, cleaved caspase-1 and IL-1β)
100 expression levels in the CCl₄+CCS group were significantly decreased (Fig. 2E-2I,
101 *P<0.05 vs. the CCl₄ group). Importantly, the expression of NLRP3, GSDMD,
102 cleaved caspase-1 and IL-1β in the CCl₄+CCS group was significantly decreased
103 compared to the expression in the CCl₄+CCS+GW4869 group, suggesting the
104 attenuating effects of Exos-rBMMSC on hepatocyte pyroptosis. In addition, IL-1β
105 and IL-18 expression levels in the CCl₄+CCS group were dramatically decreased (Fig.
106 2K-2L, ****P<0.0001 vs. the CCl₄ group). Besides, the cell counting kit-8 (CCK-8)
107 results showed that the viability of cells in the CCl₄+CCS group was significantly
108 higher than that in the other groups (Fig. 2J, ##P<0.01).

109 Exos-rBMMSCs alleviated hepatocyte pyroptosis in vitro.

110 Exos-rBMMSC were isolated by differential ultracentrifugation, as shown in Fig.
111 3A. The morphology of Exos-rBMMSC was examined by transmission electron
112 microscopy (TEM) and showed a saucer-like shape (Fig. 3B). The concentration of
113 Exos-rBMMSC was 1.0 ×10¹⁰ particles/ml, and 98% of the particles ranged from 30
114 nm to 200 nm (Fig. 3C). Exo-specific markers were highly expressed in
115 Exos-rBMMSC (Fig. 3D). Based on these results, we successfully purified
116 Exos-rBMMSC. PKH67-labelled Exos-rBMMSC were internalized by hepatocytes,
117 as shown in Fig. 3E and 3F. Exos-rBMMSC dramatically promoted CCl₄-induced
118 hepatocyte proliferation, as shown by EdU assays (Fig. 3G-3J, ****P<0.0001 vs. the
119 CCl₄ group). Similar results were obtained from the CCK-8 assay, and the cell
120 viability of the Exo group was significantly higher than that of the CCl₄ group (Fig.
121 3K, ***P<0.001). Additionally, compared to that in the CCl₄ group, IL-1β and IL-18
122 expression in the Exo group was significantly lower (Fig. 3L-3M, ***P<0.001). In
123 contrast to those in the CCl₄ group, the expression levels of pyroptosis-related
124 proteins in the Exo group were notably decreased (Fig. 3N-3R, *P<0.05).

125 The distribution of Exos-rBMMSC in vivo.

126 DiR-labelled Exos-rBMMSC were injected through the tail vein. At 0, 3 and 24
127 h postinjection, the fluorescence intensities in major organs, including the heart, lung,

128 liver, spleen and kidney, were measured by an *in vivo* imaging system. At 3 h
129 postinjection, Exos-rBMMSC were mainly concentrated in the liver and spleen. At 24
130 h postinjection, Exos-rBMMSC migrated to the lung, liver and spleen (Fig. 4A). To
131 further confirm whether PKH67-labelled Exos-rBMMSC were taken up by injured
132 hepatocytes, an immunofluorescence assay was performed. The colocalization of
133 PKH67-labelled Exos-rBMMSC with NLRP3, GSDMD and caspase-1 is shown in
134 Fig. 4B.

135 Exos-rBMMSC reduced pyroptosis protein expression *in vivo*.

136 To determine the optimal dose of Exos-rBMMSC, 100 μ g, 200 μ g and 400 μ g of
137 Exos-rBMMSC were injected via the tail vein. Four weeks postinjection, liver
138 samples from the different groups were collected. Compared to that in the CCl₄ group,
139 the expression level of NLRP3 in the 200 μ g and 400 μ g Exos-rBMMSC groups was
140 significantly decreased, while that in the 100 μ g Exos-rBMMSC group was not
141 significantly different (Fig. 4D, **P<0.01). NLRP3 expression in the 200 μ g
142 Exos-rBMMSC group was significantly lower than that in the 100 μ g Exos-rBMMSC
143 group and higher than that in the 400 μ g Exos-rBMMSC group (Fig. 4D, #P<0.05).
144 Similar results were observed for GSDMD, caspase-1 and IL-1 β loci. These proteins
145 were expressed at significantly higher levels in the CCl₄ group than in the other
146 groups (Fig. 4D-4G, **P<0.01). The expression of these proteins in the 200 μ g
147 Exos-rBMMSC group was dramatically lower than that in the 100 μ g Exos-rBMMSC
148 group but higher than that in the 400 μ g Exos-rBMMSC group (Fig. 4D-4G, #P<0.05).
149 In addition, serum IL-1 β and IL-18 levels showed similar results (Fig. 4H-4I).

150 Exos-rBMMSC further ameliorated histopathological changes.

151 HE-stained liver sections from the hepatic cirrhosis group showed inflammatory
152 cell infiltration, collagen deposition and pseudolobule formation (Fig. 5A2). Four
153 weeks postinjection, the Exos-rBMMSC-treated groups showed pathological
154 remission (Fig. 5A3-5A5). Masson-stained sections showed that 4 weeks of treatment
155 with 200 μ g and 400 μ g of Exos-rBMMSC significantly relieved collagen deposition,
156 while 100 μ g of Exos-rBMMSC induced no marked improvements compared to the
157 liver cirrhosis group (Fig. 5B, **P<0.01). Similar results were observed in the

158 semiquantitative collagen analysis; 200 μ g and 400 μ g of Exos-rBMMSC induced
159 dramatic reductions in collagen levels compared to those in the liver cirrhosis group
160 (Fig. 5K, $P<0.01$). There were no significant differences between the 100 μ g
161 Exos-rBMMSC and liver cirrhosis groups (Fig. 5K, $P>0.05$).

162 Exos-rBMMSC inhibited hepatocyte pyroptosis and accelerated proliferation.

163 Immunohistochemical assays showed that NLRP3, GSDMD and caspase-1 were
164 more highly expressed in the liver cirrhosis group than in the other groups, and the
165 levels in the 200 μ g and 400 μ g Exos-rBMMSC groups were dramatically decreased
166 (Fig. 5C-5E). Notably, proliferating cell nuclear antigen (PCNA), which is the basic
167 element in DNA replication and repair, was significantly increased in the 200 μ g and
168 400 μ g Exos-rBMMSC groups compared to the liver cirrhosis group (Fig. 5F & 5J,
169 **** $P<0.0001$). Quantitative analysis showed that NLRP3-positive cells in the 400 μ g
170 Exos-rBMMSC group were significantly reduced compared to those in the liver
171 cirrhosis group (Fig. 5G, **** $P<0.0001$) but were not significantly different compared
172 to those in the 100 μ g and 200 μ g Exos-rBMMSC groups (Fig. 5G, $P>0.05$).
173 Compared with those in the cirrhosis group, GSDMD-positive cells in the
174 Exos-rBMMSC groups were dramatically decreased (Fig. 5H, *** $P<0.001$).
175 GSDMD-positive cells in the 400 μ g Exos-rBMMSC group were markedly decreased
176 compared to those in the 200 μ g and 100 μ g Exos-rBMMSC groups (Fig. 5H,
177 ##### $P<0.0001$). Similar results were observed in the number of caspase-1-positive cells
178 (Fig. 5I).

179 Exos-rBMMSC enhanced liver function.

180 Liver function was assessed by measuring aspartate aminotransferase (AST) and
181 alanine aminotransferase (ALT) and albumin (ALB) expression levels. Compared to
182 those in the 200 μ g and 400 μ g Exos-rBMMSC groups, AST and ALT expression
183 levels in the liver cirrhosis group were markedly increased; in contrast, ALB was
184 markedly decreased (Fig. 5L-5N, * $P<0.05$). AST and ALT expression in the 200 μ g
185 Exos-rBMMSC group was significant higher than that in the 400 μ g Exos-rBMMSC
186 group, while there were no difference compared to that in the 100 μ g Exos-rBMMSC
187 group (Fig. 5L-5M, # $P<0.05$). ALB expression in the 200 μ g Exos-rBMMSC group

188 was markedly higher than that in the 100 µg Exos-rBMMSC group and dramatically
189 lower than that in the 400 µg Exos-rBMMSC group (Fig. 5N, #P<0.05).
190

191 **Discussion**

192 Liver cirrhosis is the end stage of all types of hepatic diseases, and the
193 complications cause a substantial economic burden and affect quality of life. For
194 treating liver cirrhosis and its complications, searching for effective therapies and
195 exploring the pathogenesis of liver cirrhosis have been hot topics in recent years. In
196 this study, we demonstrated that Exos-rBMMSC promoted hepatocyte proliferation
197 and alleviated liver fibrosis by restraining hepatocyte pyroptosis.

198 Pyroptosis is characterized by membrane pore formation, cytolysis and the
199 release of proinflammatory cytokines, including IL-1 β , IL-18 and HMGB1 [9].
200 Previous studies showed that hepatocyte pyroptosis and the release of inflammasome
201 components induced hepatic stellate cell (HSC) activation and liver fibrosis [10; 11].
202 Consistent with these findings, our study showed that pyroptosis-related proteins,
203 including NLRP3, caspase-1, GSDMD, cleaved caspase-1 and IL-1 β , were markedly
204 upregulated in cirrhotic liver tissues. Additionally, we confirmed that 4 weeks of
205 rBMMSC treatment by caudal vein injection significantly downregulated these
206 proteins. To further investigate the mechanism by which rBMMSCs affect
207 hepatocytes, a coculture system of rBMMSCs and hepatocytes was established, and
208 the results indicated that the effects were mediated by a paracrine pathway.

209 Exos serve as central mediators of intercellular communication owing to their
210 unique nucleic acids and proteins, which drive their utilization as cell-free
211 therapeutics [12]. Accumulating evidence has demonstrated that Exos derived from
212 stem cells alleviate liver fibrosis. The transplantation of Exos from human umbilical
213 cord-MSCs (Exos-hucMSC) mitigated liver fibrosis by reducing liver inflammation,
214 collagen deposition and epithelial-to-mesenchymal transition (EMT) [13].
215 Exos-rBMMSC could reduce oxidative stress and accelerate liver regeneration in a
216 liver injury model [14]. Additionally, Exos-rBMMSC showed suppressive effects by
217 reducing liver necrotic areas and enhancing anti-inflammatory cytokines [15]. A
218 recent study demonstrated that Exos-rBMMSC restrained liver fibrosis by inhibiting
219 HSC activation through the Wnt/ β -catenin pathway [16]. Exos produced by
220 miR-181-5p-overexpressing adipose-derived mesenchymal stem cells (ADSCs)

221 significantly inhibited collagen I, vimentin, α -SMA and fibronectin expression [17].
222 Similar research showed that circ-0000623-overexpressing ADSCs attenuated liver
223 fibrosis by activating hepatocyte autophagy [18]. However, the optimal dose of
224 Exos-rBMMSC for use in liver cirrhosis therapy has not been discussed. In this study,
225 we explored the therapeutic dose of Exos-rBMMSC for liver cirrhosis. It is worth
226 mentioning that the effect of Exos on the rat traumatic brain injury (TBI) model was
227 dose-dependent in a 7-day therapeutic window. Exos (200 μ g and 100 μ g per rat)
228 significantly alleviated long-term neuroinflammation in a TBI model, while 200 μ g
229 Exos did not provide further therapeutic effects [19]. Another study reported that
230 extracellular nanovesicles (EVs) from 1×10^8 ADSCs had better attenuating effects on
231 liver fibrosis than EVs from 1×10^6 and 1×10^7 ADSCs [20]. According to our results,
232 injection of both 200 μ g and 400 μ g of Exos-rBMMSCs via the tail vein significantly
233 restrained hepatocyte pyroptosis and collagen deposition. Interestingly, our results
234 showed that 400 μ g of Exos-rBMMSC showed a better therapeutic effect than 200 μ g
235 of Exos-rBMMSC. The present study was the first to suggest the optimal dose of
236 Exos-rBMMSCs for liver cirrhosis, laying an experimental foundation for clinical
237 applications. Further studies are warranted to discover the molecules within
238 Exos-rBMMSC, such as DNA, mRNA, noncoding RNA and specific growth factors,
239 that promote hepatocyte proliferation, restrain pyroptosis, and reduce collagen
240 deposition.
241

242 **Conclusions**

243 Overall, in this study, we demonstrated that Exos-rBMMSC significantly
244 accelerated hepatocyte proliferation and relieved liver fibrosis, and the effect may
245 depend on restraining hepatocyte pyroptosis. More importantly, the effect of
246 Exos-rBMMSC on hepatocytes was dose-dependent, and 400 µg of Exos-rBMMSC
247 may be the optimal dose for treating liver cirrhosis. Our findings provide
248 experimental data and supporting theories for clinical applications.

249

250 **Methods**

251 Establishment of the rat liver cirrhosis model

252 Sprague-Dawley rats (130~150 g, male) were obtained from the animal facility
253 of the Second Affiliated Hospital of Harbin Medical University. A liver cirrhosis
254 model was established as we described previously [3]. Briefly, 50% CCl₄ (dissolved
255 in oil) was intraperitoneally injected twice each week for 4 weeks. After 4 weeks of
256 stimulation, the animals were examined by histopathology. In addition, all
257 experiments and methods were approved by the Experimental Centre of the Second
258 Affiliated Hospital of Harbin Medical University.

259 Cell culture

260 The purification and identification of rBMSCs were described in our previous
261 study [3]. rBMSCs were cultured in DMEM/F12 medium (Sigma, USA), 10% FBS
262 (35-081-cv, Corning, USA), and 1% penicillin-streptomycin (pH = 7.4, Sigma, USA).
263 Exo-rBMSC were collected from third-passage rBMSCs to fifth-passage
264 rBMSCs. BRL rat hepatocytes were purchased from BEINA Biology Company and
265 cultured in MEM (Sigma, USA) and 10% FBS.

266 Purification and characterization of Exo-rBMSC

267 rBMSC culture supernatant was collected after the cells were cultured in
268 Exo-free medium (DMEM/F12-Sigma, 10% Exo-free FBS, 1%
269 penicillin-streptomycin) for 24 h. Then, Exos were purified according to the protocol
270 in the Journal of Extracellular Vesicles [21, 22]. Briefly, the supernatant was
271 concentrated by differential ultracentrifugation, followed by centrifugation at 300 ×g
272 for 10 min, 2000 ×g for 10 min, and 10,000 ×g for 30 min. Finally, the supernatant
273 was ultracentrifuged at 100,000 ×g for 70 min twice. The precipitate was resuspended
274 in 200 μl of phosphate-buffered saline (PBS). The protein concentration of
275 Exo-rBMSCs was measured by a BCA assay [23]. To characterize the purified
276 vesicles as Exos, the morphology was examined by TEM, and the diameters were
277 measured and analysed by a Zeta View Particle Metrix (Zeta View PMX 110). The
278 exosomal protein markers CD9, TSG101, and TSG70 (1:1000, Abcam) and negative
279 marker calnexin (1:1000, Abcam) were examined by western blotting.

280 PKH67 staining

281 According to the protocol and the published study [24], to prepare the 2× dye
282 solution, 2 μl of PKH67 ethanolic dye was added to 1 ml of Diluent C. Then, 1 ml of
283 the Exos solution was rapidly added into the 2× dye solution and mixed slowly for 4
284 min. To suspend redundant staining, 2 ml of serum-free FBS was added. PBS was
285 added to adjust the volume to 14 ml, and the mixture was centrifuged at 100,000 × g
286 at 4 °C for 90 min. The supernatant was removed, and PKH67-labelled Exos were
287 resuspended in 200 μl of PBS. Similarly, rBMMSCs were stained with PKH67
288 according to the Exo procedure.

289 Coculture of hepatocytes and rBMMSC

290 The Transwell coculture system (3450, Corning Costar) consisted of 1×10⁶
291 hepatocytes and 1×10⁶ rBMMSCs. Hepatocytes were seeded in 6-well plates, and
292 PKH67-labelled rBMMSCs were cultured in the Transwell inserts. After 24 h of
293 coculture, the hepatocytes were observed under a fluorescence microscope
294 (Zeiss-DMI8).

295 Exo uptake assessment

296 PKH67-labelled Exos-rBMMSC (100 μg) were added to hepatocytes and
297 cultured in 6-well plates with FBS-free medium. After 24 h of incubation, the
298 hepatocytes were assessed by fluorescence microscopy.

299 Cell viability analysis

300 The viability of untreated hepatocytes, hepatocytes treated with CCl₄ (10 mM)
301 and PBS, and hepatocytes treated with CCl₄ (10 mM) and 100 μg Exos was assessed
302 by CCK-8 (APExBIO-K1018) and 5-ethynyl-2'-deoxyuridine (EdU) imaging kits (UE,
303 China). Briefly, 5×10³ hepatocytes were seeded in 96-well plates for 24 h (n = 15).
304 Then, CCl₄ (10 mM) was added to 10 plates, and the same volume of PBS was added
305 to 5 plates. After 12 h of incubation, the medium was replaced with MEM+100 μg
306 Exos or MEM+PBS. Cell viability was measured according to the manufacturer's
307 protocol after 4 hours of culture.

308 Biodistribution of Exos-rBMMSC

309 To detect the distribution of the Exos-rBMMSC in vivo, Exos-rBMMSC were

310 stained with 1,1'-dioctadecyl-3,3,3',3'-tetramethylindotricarbocyanine iodide (DiR,
311 D12731, Invitrogen, Life Technologies), which was obtained commercially from
312 Sigma. DiR-stained Exos (100 µg) were injected into CCl₄-induced rats. At 0, 3 and
313 24 h after tail vein injection, ex vivo fluorescent images of different organs, including
314 the heart, lung, liver, spleen and kidney, were taken by an in vivo imaging system
315 (Night OWL II LB983).

316 Immunofluorescence and immunohistochemical assays

317 Immunofluorescence and immunohistochemical examinations were conducted as
318 described previously [3]. Briefly, frozen liver sections (4 µm) were used for
319 immunofluorescence assays, and paraffin liver sections (4 µm) were used for
320 immunohistochemical assays. Primary antibodies against NLRP3 (1:50, NOVUS),
321 GSDMD (1:1000, Abcam), Caspase-1 (1:200, Sigma), and PCNA (1:200, Affinity)
322 were added and incubated overnight at 4 °C. Then, immunofluorescence images were
323 taken by fluorescence microscopy (Zeiss-DMI8), and immunohistochemical images
324 were obtained by an Olympus (BX41) microscope and semiquantitatively measured
325 by Fiji software.

326 Western blotting

327 The western blot procedure was described in our previous study [3]. Briefly,
328 proteins were extracted from frozen liver tissue. After electrophoresis, the proteins
329 were transferred to nitrocellulose membranes, which were incubated with β-actin
330 (1:1000, Abcam), NLRP3 (1:5000, NOVUS), GSDMD (1:1000, Abcam), cleaved
331 caspase-1 (1:1000, CST), and IL-1β (1:1000, Abclonal) antibodies overnight at 4 °C.
332 Then, the nitrocellulose membranes were incubated with HRP-conjugated secondary
333 antibodies (1:5000, Boster). Finally, images were taken using the ImageQuant LAS
334 4000 mini machine (GE).

335 ELISA

336 Rat tissue and serum samples were obtained after euthanasia. Fresh frozen
337 hepatic tissues were homogenized with PBS at 4 °C according to the manufacturer's
338 protocol. After centrifugation at 120,000 g at 4 °C for 20 min, the supernatant was
339 collected. Liver function was evaluated by measuring the levels of ALT, AST and

340 ALB with ELISA kits (Nanjing Jiancheng, China). The IL-1 β and IL-18 levels in
341 serum and cell supernatants were examined by ELISA kits (Boster, China) according
342 to the protocol.

343 Histopathological assays

344 The liver tissues were fixed with 4% paraformaldehyde and embedded in
345 paraffin. Liver sections (4 μ m) were stained with HE to observe pathological
346 morphology and with Masson to examine collagen deposition. Semiquantitative
347 analysis was performed by Fiji software.

348 Statistical analyses

349 GraphPad Prism 9 software (Version 9.0.1, Inc.) was used to analyse the data by
350 one-way analysis of variance (ANOVA). All data are displayed as the mean \pm
351 standard deviation. Statistical significance was defined as $P < 0.05$. All figures were
352 prepared by GraphPad Prism 9 and Photoshop 7.1.

353

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427

428 **Figure legends**

429 **Figure 1. The alleviation of pyroptosis by rBMMSCs in the context of liver**

430 **cirrhosis.** Specimens of the control and cirrhotic liver models are shown in Fig. 1A.

431 Pathological HE- and Masson-stained sections of control and cirrhotic liver tissues are

432 shown (Fig. 1B-1E). Pyroptosis-related proteins were highly expressed in the liver

433 cirrhosis model (Fig. 1F). Pathological sections showed significant alleviation in the

434 rBMMSC transplantation group (Fig. 1G-1H). Pyroptosis-related proteins in the

435 BMMSC group were significantly downregulated compared to those in the liver

436 cirrhosis group (Fig. 1I-1M, * $P < 0.05$ vs. the CCl_4 group). IL-1 β and IL-18 expression

437 in the BMMSC group was dramatically decreased (Fig. 1N-1O, ** $P < 0.01$ vs. the CCl_4

438 group). The data are displayed as the means \pm SD.

439 **Figure 2. Paracrine effect by which rBMMSCs inhibit hepatocyte pyroptosis.**

440 PKH67 dye was transferred to hepatocytes by rBMMSCs via the paracrine pathway

441 (Fig. 2A-2D). Pyroptosis-related proteins in the CCl_4 +CCS group were significantly

442 decreased compared to those in the CCl_4 group and CCl_4 +CCS+GW4869 group (Fig.

443 2E-2I, ** $P < 0.01$). The expression of pyroptosis-related proteins in the CCl_4 +DMSO

444 and CCl_4 +GW4869 groups was not significantly different compared to that in the

445 CCl_4 group (Fig. 2E-2I). Cell viability in the CCl_4 +CCS group was significantly

446 higher than that in the other groups (Fig. 2J, ## $P < 0.01$). The expression of IL-1 β and

447 IL-18 in the CCl_4 +CCS group was dramatically lower than that in the other treatment

448 groups (Fig. 2K-2L, ** $P < 0.01$). The data are displayed as the means \pm SD.

449 **Figure 3. Effects of Exos-rBMMSC on CCl_4 -induced hepatocytes.** The extraction

450 protocol of Exos-rBMMSCs is shown (Fig. 3A). TEM morphology and nanoparticle

451 tracking analysis (NTA) results are shown (Fig. 3B-3C). HSP70, TSG101, CD9 and

452 calnexin expression in the rBMMSC and Exos-rBMMSC groups is shown (Fig. 3D).

453 PKH67-labelled Exos-rBMMSC were internalized by hepatocytes, as observed under

454 a fluorescence microscope (Fig. 3E-3F). Hepatocyte proliferation in the CCl_4 group

455 was dramatically increased (Fig. 3G-3K, *** $P < 0.001$ vs. the CCl_4 group). IL-1 β and

456 IL-18 expression in the Exo group was significantly decreased (Fig. 3L-3M,

457 *** $P < 0.001$ vs. the CCl_4 group). NLRP3, GSDMD, cleaved caspase-1 and IL-1 β

458 expression in the Exo group was significantly decreased compared to that in the CCl₄
459 group (Fig. 3N-3R, *P<0.05). The data are displayed as the means ± SD.

460 **Figure 4. Distribution of Exos-rBMMSC and the therapeutic effect on liver**
461 **cirrhosis.** The distribution of Exos-rBMMSCs was examined by NIRF at 0 h, 3 h and
462 24 h postinjection. At 3 h postinjection, Exos-rBMMSC were mainly concentrated in
463 the liver and were mainly concentrated in the liver and lung at 24 h postinjection (Fig.
464 4A). The colocalization of NLRP3, GSDMD and caspase-1 with PKH67 dye are
465 shown in Fig. 4B. Compared to that in the CCl₄ group, NLRP3, GSDMD, cleaved
466 caspase-1, and IL-1β expression in the 200 μg and 400 μg Exos-rBMMSC groups was
467 dramatically decreased (Fig. 4C-4G, **P<0.01). The expression of NLRP3 in the 100
468 μg Exos-rBMMSC group was not significantly different compared to that in the CCl₄
469 group (Fig. 4D, P>0.05). The expression of GSDMD, cleaved caspase-1 and IL-1β in
470 the 100 μg Exos-rBMMSC group was dramatically decreased compare to that in the
471 CCl₄ group (Fig. 4E-4G, **P<0.01). The expression of NLRP3, GSDMD, cleaved
472 caspase-1 and IL-1β in the 200 μg Exos-rBMMSC group was significantly higher
473 than that in the 400 μg Exos-rBMMSC group (Fig. 4C-4G, #P<0.05). IL-1β and IL-18
474 expression in the 200 μg and 400 μg Exos-rBMMSC groups were dramatically lower
475 than that in the CCl₄ group (Fig. 4H-4I, ###P<0.001), while the expression in the 100
476 μg Exos-rBMMSC group was not different from that in the CCl₄ group (Fig. 4H-4I,
477 P>0.05). IL-1β and IL-18 expression in the 200 μg Exos-rBMMSC group were
478 significantly lower than that in the 100 μg Exos-rBMMSC group but was higher than
479 that in the 400 μg Exos-rBMMSC group (Fig. 4H-4I, ##P<0.01). The data are
480 displayed as the means ± SD.

481 **Figure 5. The therapeutic effect of Exos-BMMSC on liver cirrhosis was assessed**
482 **in pathological sections and by measuring liver function.** HE-stained sections from
483 the control, liver cirrhosis and Exos-rBMMSC-treated groups are shown (Fig. 5A).
484 Masson-stained sections from the different treatment groups are shown (Fig. 5B).
485 Semiquantitative analysis of the collagen-positive area showed that the areas in the
486 200 μg and 400 μg Exos-rBMMSC groups were significantly decreased compared
487 with those in the liver cirrhosis group (Fig. 5K, **P<0.01). The collagen area in the

488 200 µg Exos-rBMMSC group was significantly lower than that in the 100 µg
489 Exos-rBMMSC group but was higher than that in the 400 µg Exos-rBMMSC group
490 (Fig. 5K, #P<0.05). NLRP3-, GSDMD-, caspase-1- and PCNA-positive cells were
491 assessed by immunohistochemistry (Fig. 5C-5F). Semiquantitative analysis showed
492 that NLRP3-, GSDMD-, caspase-1- and PCNA-positive cells in the 100 µg, 200 µg
493 and 400 µg Exos-rBMMSC groups were significantly lower than those in the liver
494 cirrhosis group (Fig. 5G-5J, **P<0.01). The numbers of GSDMD-, caspase-1- and
495 PCNA-positive cells in the 200 µg Exos-rBMMSC group were dramatically lower
496 than those in the 100 µg Exos-rBMMSC group but were higher than those in the 400
497 µg Exos-rBMMSC group (Fig. 5G-5J, #P<0.05). NLRP3-positive cells in the 200 µg
498 Exos-rBMMSC group were not significantly different from those in the 100 µg and
499 400 µg Exos-rBMMSC groups (Fig. 5G). AST and ALT expression in the 200 µg and
500 400 µg Exos-rBMMSC groups was significantly decreased, but ALB expression was
501 significantly increased compared to that in the liver cirrhosis group (Fig. 5L-5N,
502 *P<0.05). The expression of AST and ALT in the 400 µg Exos-rBMMSC group was
503 dramatically lower, while the expression of ALB was higher than that in the 200 µg
504 Exos-rBMMSC group (5L-5N, #P<0.05). The data are displayed as the means ± SD.
505

506 **Abbreviations**

507 Exos: Exosomes; rBMMSC: rat bone marrow mesenchymal stem cells;
508 Exos-rBMMSC: Exosomes derived from rat bone marrow mesenchymal stem cell;
509 CCS: cell culture supernatant; EVs: Extracellular nanovesicles; miRNAs: mirco-RNA;
510 Exos-hucMSC: Exos from human umbilical cord-MSCs; NIRF: Near-infrared
511 fluorescence; HE: Hematoxylin and Eosin; ADSCs:Adipose-derived mesenchymal
512 stem cells; EMT: Epithelial-to-mesenchymal transition; TBI: Traumatic brain injury;
513 TEM: Transmission electronic microscopy; CCK8: Cytotoxicity kit-8; EdU:
514 5-Ethynyl-2'-deoxyuridine; AST: aspartate aminotransferase; DiR:
515 1,1'-Dioctadecyl-3,3,3',3'-tetramethylindotricarbocyanine iodide; ALT: alanine
516 aminotransferase; ALB: albumin; ANOVA: One-way analysis of variance.

517 **Authors' contributions**

518 Zhang YC designed and completed experiments. Jin SZ and Nie XS involved in
519 experiment design. Zhang YC, Nie XS and Wang LJ contributed to paper writing. Jin
520 H, Pu RH and Liang MH contributed to data analysis. Chang Y, Gao Y and Zhang HL
521 involved in drawing scientific pictures. All authors read and approved the final
522 manuscript.

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527 **Availability of data and materials**

528 All data generated or analysed during this study are included in this published
529 article.

530 **Consent for publication**

531 All authors agree to be published.

532 **Competing interests**

533 All authors declare no conflicts of interest in this work.

534 **Ethics approval and consent to participate**

535 We confirmed all experiments and methods performed in this study in

536 accordance with the 2000 Helsinki Declaration and were approved by the
537 Experimental Center of the Second Affiliated Hospital of Harbin Medical University.

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Figures

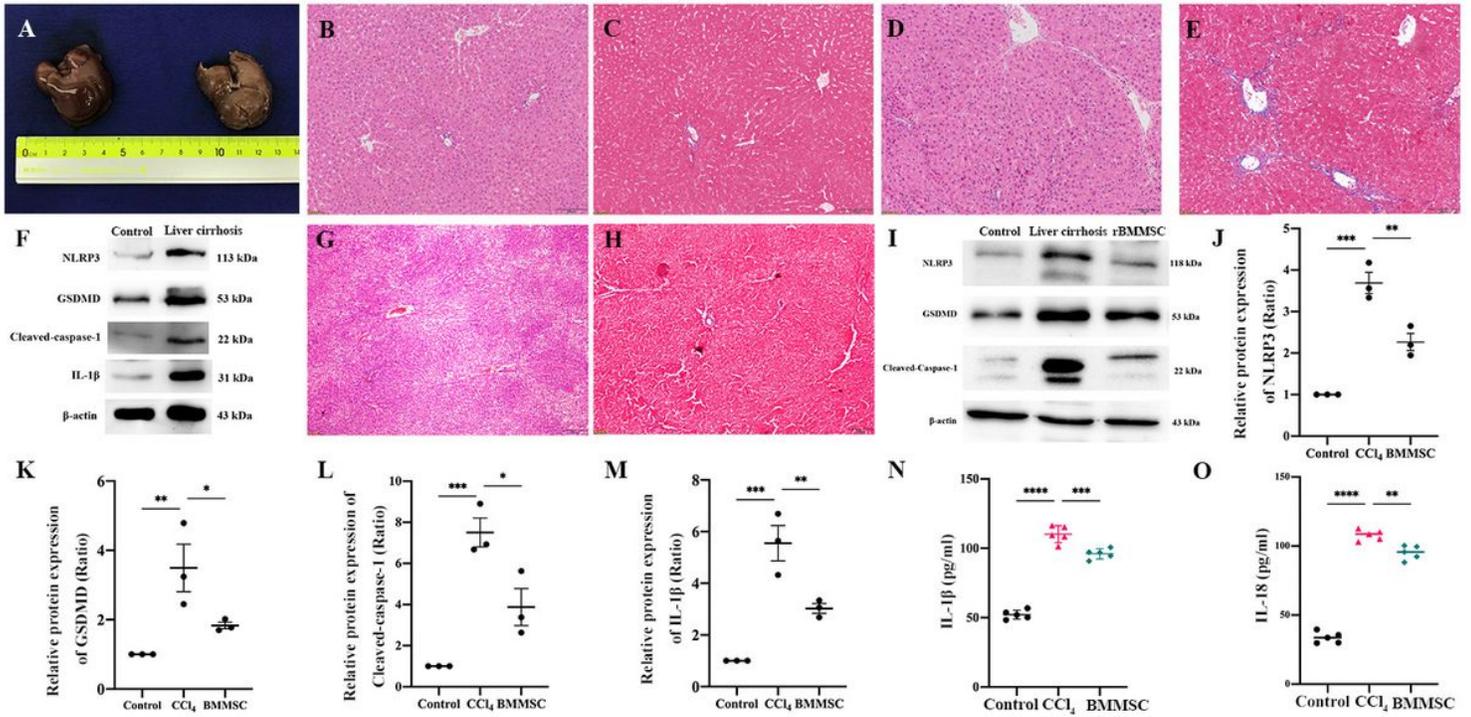


Figure 1

The alleviation of pyroptosis by rBMMSCs in the context of liver cirrhosis. Specimens of the control and cirrhotic liver models are shown in Fig. 1A. Pathological HE- and Masson-stained sections of control and cirrhotic liver tissues are shown (Fig. 1B-E). Pyroptosis-related proteins were highly expressed in the liver cirrhosis model (Fig. 1F). Pathological sections showed significant alleviation in the rBMMSC transplantation group (Fig. 1G-H). Pyroptosis-related proteins in the BMMSC group were significantly downregulated compared to those in the liver cirrhosis group (Fig. 1I-M, * $P < 0.05$ vs. the CCl₄ group). IL-1 β and IL-18 expression in the BMMSC group was dramatically decreased (Fig. 1N-O, ** $P < 0.01$ vs. the CCl₄ group). The data are displayed as the means \pm SD.

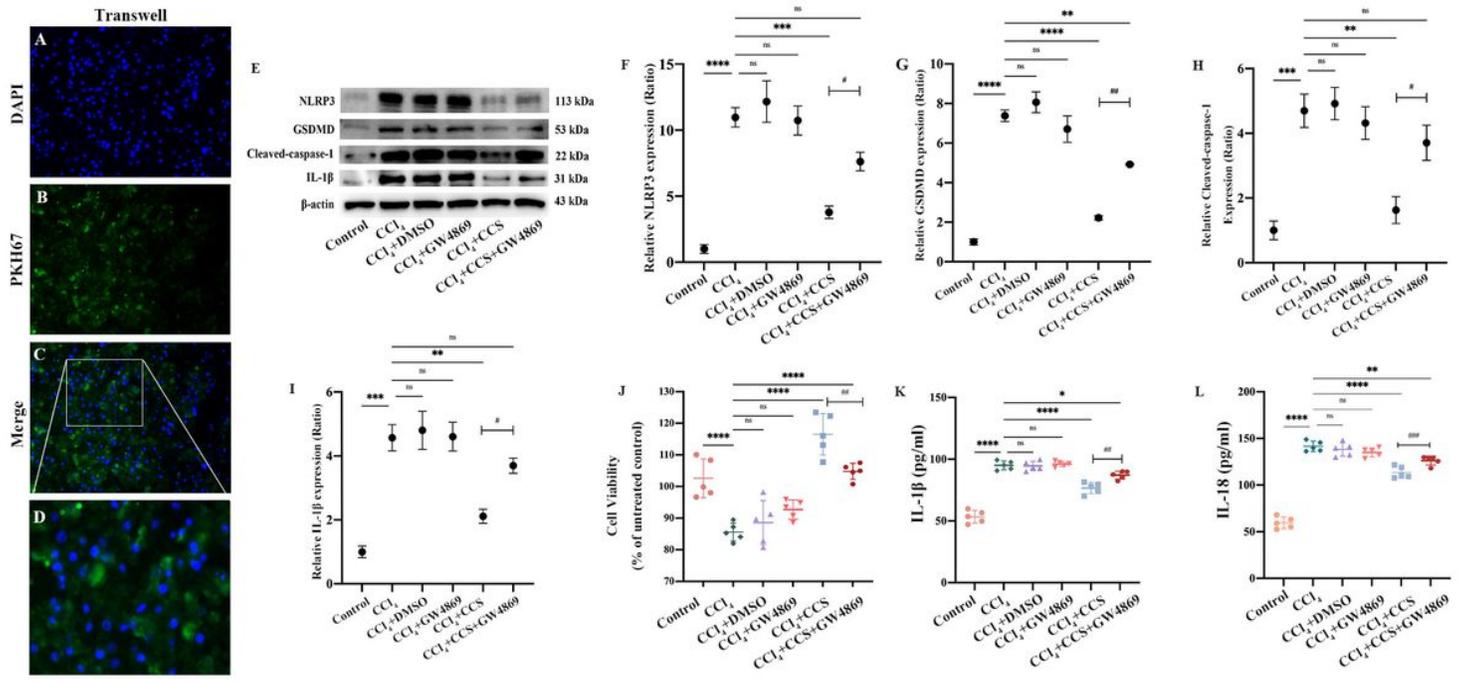


Figure 2

Paracrine effect by which rBMMSCs inhibit hepatocyte pyroptosis. PKH67 dye was transferred to hepatocytes by rBMMSCs via the paracrine pathway (Fig. 2A-2D). Pyroptosis-related proteins in the CCl₄+CCS group were significantly decreased compared to those in the CCl₄ group and CCl₄+CCS+GW4869 group (Fig. 2E-2I, **P<0.01). The expression of pyroptosis-related proteins in the CCl₄+DMSO and CCl₄+GW4869 groups was not significantly different compared to that in the CCl₄ group (Fig. 2E-2I). Cell viability in the CCl₄+CCS group was significantly higher than that in the other groups (Fig. 2J, ##P<0.01). The expression of IL-1β and IL-18 in the CCl₄+CCS group was dramatically lower than that in the other treatment groups (Fig. 2K-2L, **P<0.01). The data are displayed as the means ± SD.

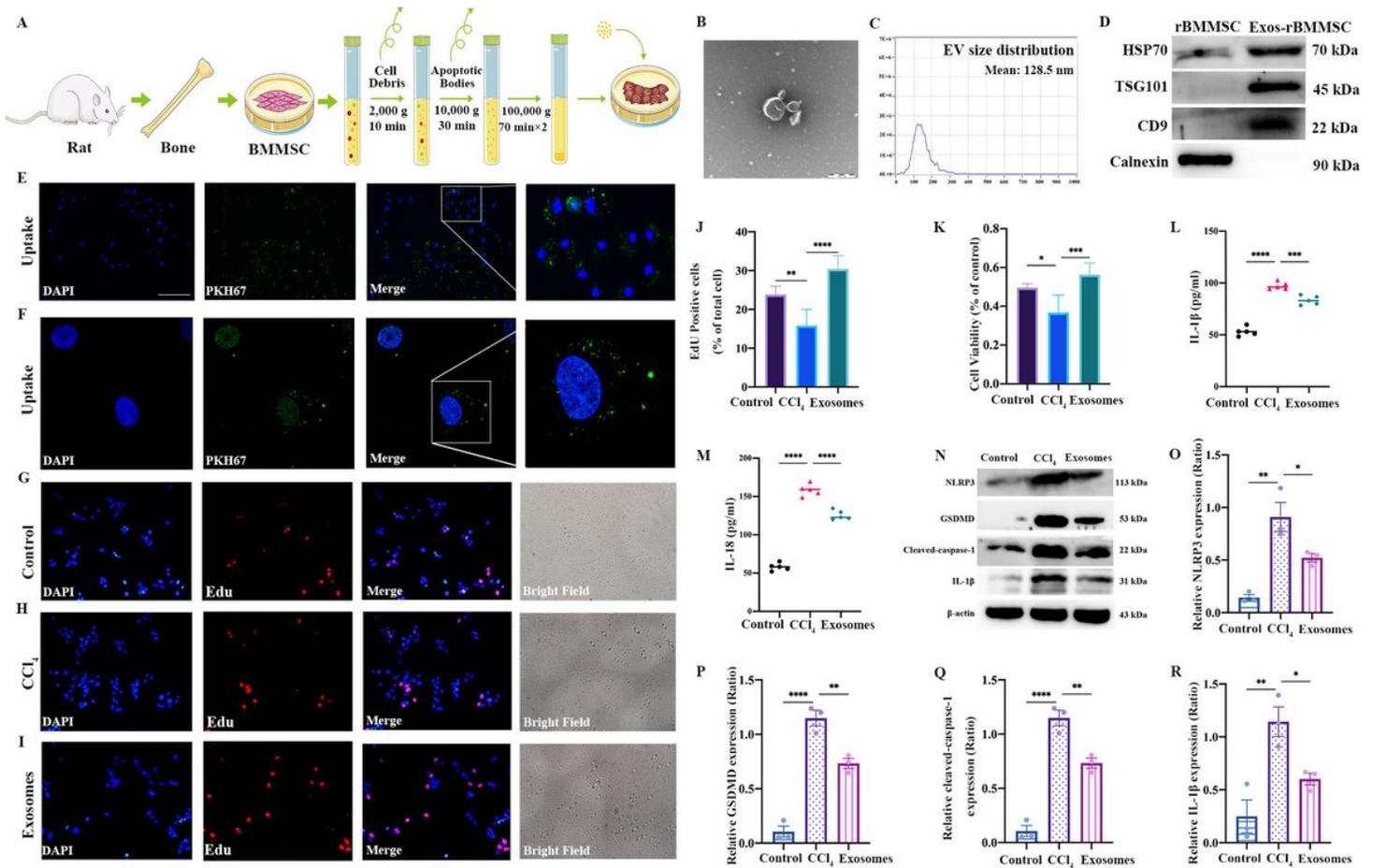


Figure 3

Effects of Exos-rBMMSC on CCl₄-induced hepatocytes. The extraction protocol of Exos-rBMMSCs is shown (Fig. 3A). TEM morphology and nanoparticle tracking analysis (NTA) results are shown (Fig. 3B-3C). HSP70, TSG101, CD9 and calnexin expression in the rBMMSC and Exos-rBMMSC groups is shown (Fig. 3D). PKH67-labelled Exos-rBMMSC were internalized by hepatocytes, as observed under a fluorescence microscope (Fig. 3E-3F). Hepatocyte proliferation in the CCl₄ group was dramatically increased (Fig. 3G-3K, ****P*<0.001 vs. the CCl₄ group). IL-1β and IL-18 expression in the Exo group was significantly decreased (Fig. 3L-3M, ****P*<0.001 vs. the CCl₄ group). NLRP3, GSDMD, cleaved caspase-1 and IL-1β expression in the Exo group was significantly decreased compared to that in the CCl₄ group (Fig. 3N-3R, **P*<0.05). The data are displayed as the means ± SD.

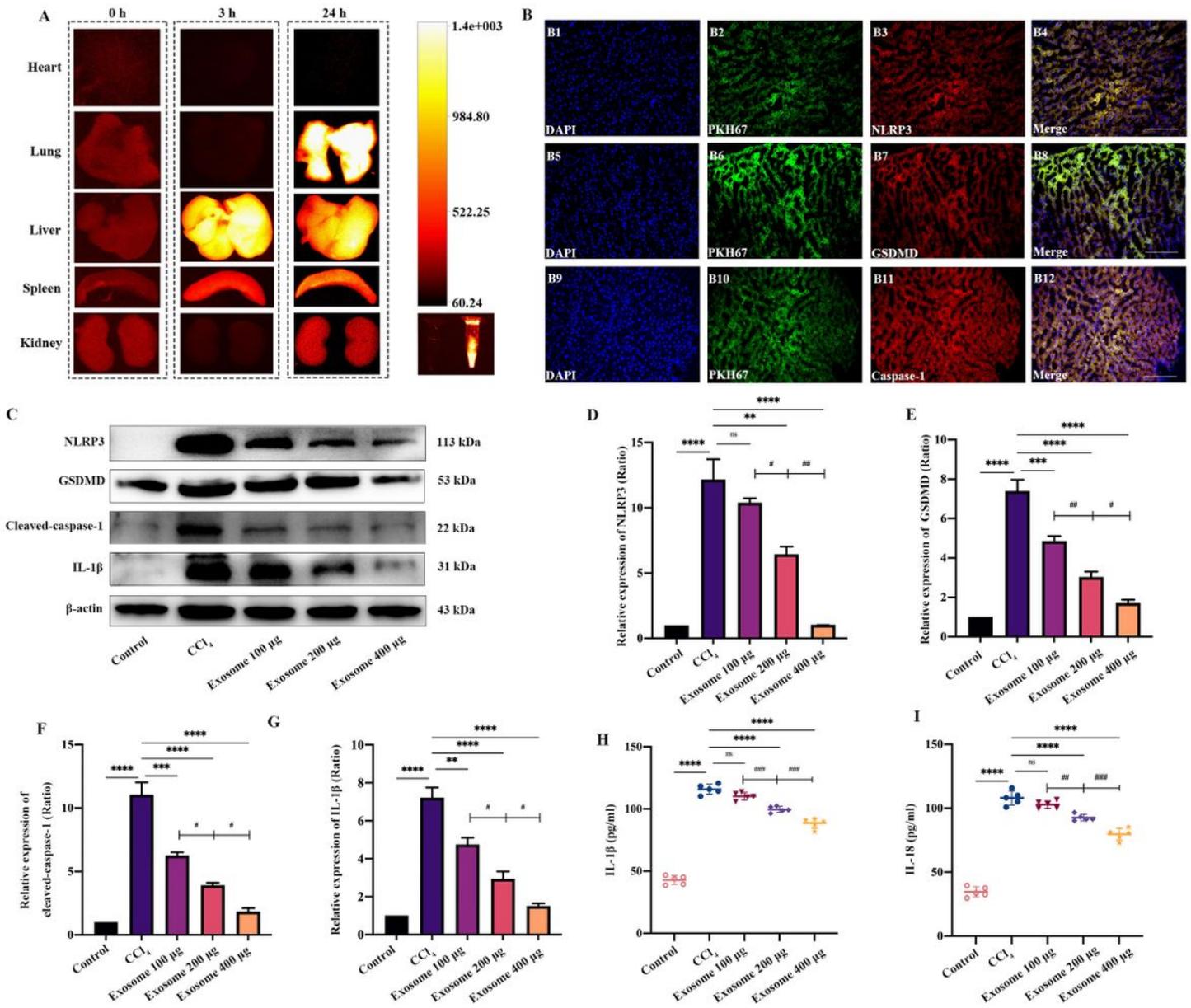


Figure 4

Distribution of Exos-rBMMSC and the therapeutic effect on liver cirrhosis. The distribution of Exos-rBMMSCs was examined by NIRF at 0 h, 3 h and 24 h postinjection. At 3 h postinjection, Exos-rBMMSC were mainly concentrated in the liver and were mainly concentrated in the liver and lung at 24 h postinjection (Fig. 4A). The colocalization of NLRP3, GSDMD and caspase-1 with PKH67 dye are shown in Fig. 4B. Compared to that in the CCl₄ group, NLRP3, GSDMD, cleaved caspase-1, and IL-1β expression in the 200 μg and 400 μg Exos-rBMMSC groups was dramatically decreased (Fig. 4C-4G, **P<0.01). The expression of NLRP3 in the 100 μg Exos-rBMMSC group was not significantly different compared to that in the CCl₄ group (Fig. 4D, P>0.05). The expression of GSDMD, cleaved caspase-1 and IL-1β in the 100 μg Exos-rBMMSC group was dramatically decreased compare to that in the CCl₄ group (Fig. 4E-4G, **P<0.01). The expression of NLRP3, GSDMD, cleaved caspase-1 and IL-1β in the 200 μg Exos-rBMMSC group was significantly higher than that in the 400 μg Exos-rBMMSC group (Fig. 4C-4G, #P<0.05). IL-1β

and IL-18 expression in the 200 μ g and 400 μ g Exos-rBMMSC groups were dramatically lower than that in the CCl₄ group (Fig. 4H-4I, ## P <0.001), while the expression in the 100 μ g Exos-rBMMSC group was not different from that in the CCl₄ group (Fig. 4H-4I, P >0.05). IL-1 β and IL-18 expression in the 200 μ g Exos-rBMMSC group were significantly lower than that in the 100 μ g Exos-rBMMSC group but was higher than that in the 400 μ g Exos-rBMMSC group (Fig. 4H-4I, ## P <0.01). The data are displayed as the means \pm SD.

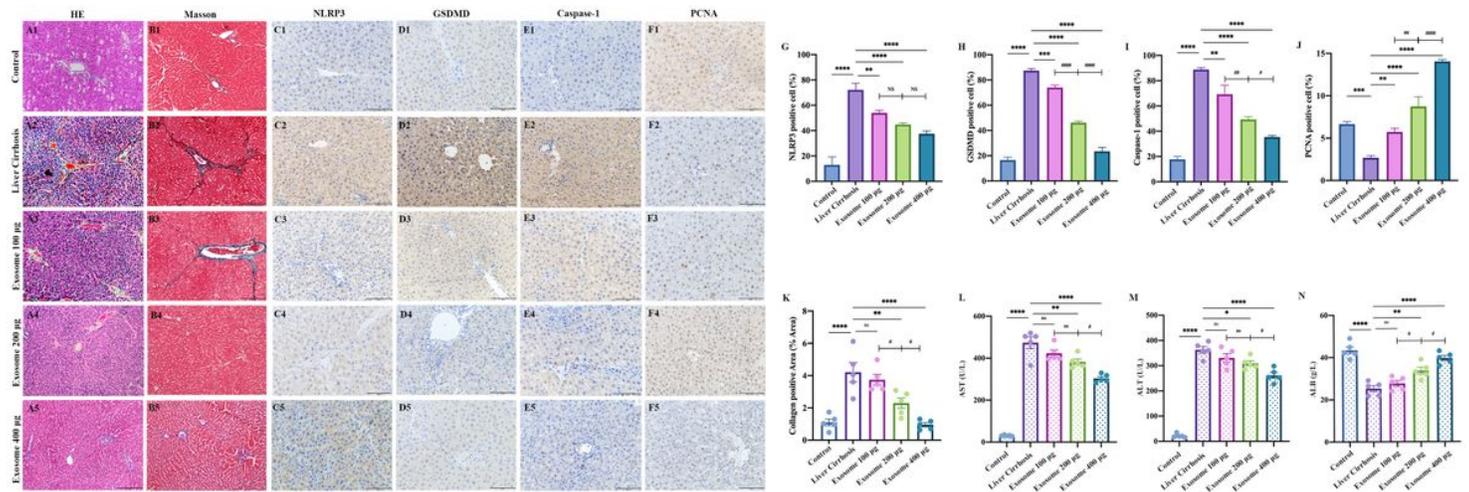


Figure 5

The therapeutic effect of Exos-BMMSC on liver cirrhosis was assessed in pathological sections and by measuring liver function. HE-stained sections from the control, liver cirrhosis and Exos-rBMMSC-treated groups are shown (Fig. 5A). Masson-stained sections from the different treatment groups are shown (Fig. 5B). Semiquantitative analysis of the collagen-positive area showed that the areas in the 200 μ g and 400 μ g Exos-rBMMSC groups were significantly decreased compared with those in the liver cirrhosis group (Fig. 5K, ** P <0.01). The collagen area in the 200 μ g Exos-rBMMSC group was significantly lower than that in the 100 μ g Exos-rBMMSC group but was higher than that in the 400 μ g Exos-rBMMSC group (Fig. 5K, # P <0.05). NLRP3-, GSDMD-, caspase-1- and PCNA-positive cells were assessed by immunohistochemistry (Fig. 5C-5F). Semiquantitative analysis showed that NLRP3-, GSDMD-, caspase-1- and PCNA-positive cells in the 100 μ g, 200 μ g and 400 μ g Exos-rBMMSC groups were significantly lower than those in the liver cirrhosis group (Fig. 5G-5J, ** P <0.01). The numbers of GSDMD-, caspase-1- and PCNA-positive cells in the 200 μ g Exos-rBMMSC group were dramatically lower than those in the 100 μ g Exos-rBMMSC group but were higher than those in the 400 μ g Exos-rBMMSC group (Fig. 5G-5J, # P <0.05). NLRP3-positive cells in the 200 μ g Exos-rBMMSC group were not significantly different from those in the 100 μ g and 400 μ g Exos-rBMMSC groups (Fig. 5G). AST and ALT expression in the 200 μ g and 400 μ g Exos-rBMMSC groups was significantly decreased, but ALB expression was significantly increased compared to that in the liver cirrhosis group (Fig. 5L-5N, * P <0.05). The expression of AST and ALT in the 400 μ g Exos-rBMMSC group was dramatically lower, while the expression of ALB was higher than that in the 200 μ g Exos-rBMMSC group (5L-5N, # P <0.05). The data are displayed as the means \pm SD.

Supplementary Files

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