ORIGINAL ARTICLE

# MCP1 triggers monocyte dysfunctions during abnormal osteogenic differentiation of mesenchymal stem cells in ankylosing spondylitis

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## Abstract

Ankylosing spondylitis (AS) is a chronic inflammatory disease characterized by pathological osteogenesis and inflammation. However, the pathogenesis of AS and the pathological relationship between osteogenesis and inflammation in this disease remain largely unknown. Mesenchymal stem cells (MSCs) are multipotent progenitor cells capable of osteogenic differentiation and immunoregulation. Recently, we demonstrated that MSCs from AS patients (ASMSCs) have a greater potential for osteogenic differentiation than MSCs from healthy donors (HDMSCs), which therefore seems to be a component of pathological osteogenesis in AS. Previous studies have indicated that the immunoregulatory abilities of MSCs change following differentiation. However, the subsequent effects of ASMSCs during abnormal osteogenic differentiation are unclear. Here, we further demonstrated that ASMSCs secreted more monocyte chemoattractant protein 1 (MCP1) than HDMSCs during osteogenic differentiation.

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This enhanced MCP1 secretion augmented monocyte migration, increased classical macrophage polarization, and enhanced TNF- $\alpha$  secretion. Inhibiting MCP1 secretion from osteogenic differentiated ASMSCs using lentiviruses encoding short hairpin RNAs ameliorated these dysfunctions. Blocking the ERK1/2 pathway in ASMSCs with U0126 corrected the abnormal osteogenic differentiation, inhibited MCP1 overexpression, and prevented subsequent monocyte dysfunction. Finally, MCP1 expression was up-regulated during osteogenic differentiation in ASMSCs in vivo and was locally augmented in osteoblasts at ossification sites in AS patients. In summary, our study determined that MCP1 overexpression during abnormal osteogenic differentiation of ASMSCs triggers monocyte dysfunctions. We propose the novel hypothesis that pathological osteogenesis can lead to inflammation in AS. This hypothesis may contribute to reveal the precise pathological relationship between osteogenesis and inflammation in the field of osteoimmunology.

#### Key message

- & ASMSCs secreted more MCP1 during abnormal osteogenic differentiation.
- MCP1 overexpression leads to monocyte dysfunctions.
- Pathological osteogenesis can lead to inflammation in AS.

Keywords Ankylosing spondylitis · Mesenchymal stem cells . Monocyte chemoattractant protein 1 . Monocyte

## Introduction

Ankylosing spondylitis (AS) is a chronic inflammatory disease characterized by chronic inflammation and pathological osteogenesis [\[1](#page-10-0)]. Increasing research efforts have focused on the pathogenesis of AS. However, the precise mechanisms underlying inflammation and pathological osteogenesis remain poorly understood, and controversies exist regarding the relationship between these two processes in AS [\[2,](#page-10-0) [3](#page-10-0)]. Some studies have demonstrated that inflammation predominates and results in pathological osteogenesis [\[4,](#page-10-0) [5](#page-10-0)]. Other studies have shown that these two processes are uncoupled [\[6,](#page-10-0) [7\]](#page-10-0). However, whether pathological osteogenesis leads to inflammation in AS has never been investigated.

Mesenchymal stem cells (MSCs), multipotent stromal cells capable of immunoregulation and tri-lineage differentiation, participate in both inflammation and osteogenesis [\[8,](#page-10-0) [9](#page-10-0)]. It is acknowledged that MSC dysfunctions contribute to various inflammatory diseases [[10](#page-10-0)–[12](#page-10-0)]. We recently demonstrated that because they secrete more bone morphogenetic protein 2 (BMP2) but less Noggin, MSCs from AS patients (ASMSCs) have greater capacities for osteogenic differentiation than MSCs from healthy donors (HDMSCs), which is a mechanism underlying pathological osteogenesis in AS [\[13](#page-10-0)]. It has been reported that immunoregulatory abilities of MSCs change following differentiation [\[14\]](#page-10-0). Whether ASMSCs lead to subsequent dysfunctions, especially in immunoregulation, during abnormal osteogenic differentiation remains unclear.

Chemokines, immunomodulatory factors secreted by MSCs, are small heparin-binding proteins that participate in inflammatory reactions [[15](#page-10-0)]. Monocyte chemoattractant protein 1 (MCP1), known as C-C motif chemokine ligand 2 (CCL2), is one of the most important chemokines [\[15](#page-10-0), [16\]](#page-10-0). Produced by various types of cells including MSCs, MCP1 affects monocyte migration and subsequent macrophage polarization [\[16,](#page-10-0) [17\]](#page-10-0). Previous studies have demonstrated that MCP1 parasecretion contributes to the development of inflammatory diseases, including AS [[18](#page-10-0)–[22](#page-10-0)]. Particularly, MCP1 oversecretion during MSC osteogenic differentiation induces immune disorder [\[23](#page-10-0)]. However, MCP1 expression patterns and their effects on monocyte during abnormal osteogenic differentiation of ASMSCs remain unknown.

Here, we demonstrated that because of extracellular signalregulated kinase 1/2 (ERK1/2) pathway overactivation, ASMSCs secreted more MCP1 during abnormal osteogenic differentiation than HDMSCs, resulting in augmented monocyte migration, enhanced M1 macrophage polarization, and increased tumor necrosis factor-α (TNF-α) secretion. We speculate that MCP1 triggers monocyte dysfunction during abnormal osteogenic differentiation of ASMSCs and propose the novel hypothesis that pathological osteogenesis leads to inflammation in AS.

# Materials and methods

#### Ethics statement

Welfare Committee of Sun Yat-Sen Memorial Hospital, Sun Yat-Sen University, Guangzhou, China. Before the study, all healthy donors and AS patients were informed of the clinical requirements and possible risks of all operations, and signed informed consents were obtained.

# Cell isolation and culture

Twenty-one healthy donors and 19 AS patients were recruited. AS patients were diagnosed according to the modified New York criteria [\[24](#page-11-0)]. The characteristics of study subjects are presented in Supplemental Table 1. After bone marrow punctures, MSCs were immediately isolated and cultured as de-scribed [[13\]](#page-10-0). MSCs were used at passage 3.

Peripheral blood mononuclear cells (PBMCs) were isolated via density gradient centrifugation. CD14<sup>+</sup> monocytes were isolated from PBMCs using CD14 MicroBeads (Miltenyi Biotec).

#### Flow cytometry

To assess the purity,  $CD14<sup>+</sup>$  monocytes were incubated with CD14 antibody (Miltenyi Biotec). For macrophage polarization assays, cells were incubated with antibodies against HLA-DR-PE or CD206-PE (BD Pharmingen) and then incubated with fixation medium (Invitrogen) for 15 min. After three washes, cells were incubated with a permeabilization medium (Invitrogen) plus CD68 antibody (BD Pharmingen). All labeled cells were detected using BD Influx cell sorter (BD Biosciences).

### Osteogenic differentiation

MSCs were seeded in 12-well plates at a density of  $1.5 \times 10^4$  cells/cm<sup>2</sup> and cultured in osteogenic differentiation medium composed of Dulbecco's modified Eagle's medium (DMEM, GIBCO) supplemented with 10% fetal bovine serum (FBS, GIBCO), 0.1 μM dexamethasone, 10 mM βglycerol phosphate, 50 μM ascorbic acid, 100 IU/ml penicillin, and 100 IU/ml streptomycin (Sigma-Aldrich). The culture medium was replaced every 3 days.

# Quantitative real-time PCR

Total RNAwas extracted using RNAiso Plus and reserve transcribed into cDNA using PrimeScript™ reagent kits (TaKaRa). High-throughput microfluidics fluorescence qRT-PCR was performed as described [\[25\]](#page-11-0). Briefly, sample premix and assay pre-mix were added into a 48.48 Dynamic Array Integrated Fluidic Circuit (IFC; Fluidigm), which was primed and loaded in IFC Controller (Fluidigm). Reactions were performed in BioMark™ HD System (Fluidigm). Data were first normalized to GAPDH expression and second to 0 day values of HDMSCs. Data were analyzed with PCR analysis software (Fluidigm). For ordinary qRT-PCR, reactions were performed in the LightCycler®480 PCR system (Roche) using SYBR® Premix Taq™ kits (TaKaRa). The relative expression levels were analyzed using the  $2^{-\Delta\Delta Ct}$ method. Primers for each gene are shown in Supplemental Table 2.

#### Migration assay

Migration assays were performed using Polycarbonate Membrane Transwell® Inserts (5.0-μm pores; Corning). MSCs ( $2 \times 10^4$ ) in 600 µl culture supernatant were placed in the lower chambers. In some assays, 600 μl cell-free MSC culture supernatant was placed in the lower chambers. After 8 h, the upper chambers, containing  $1 \times 10^6$  CD14<sup>+</sup> monocytes suspended in 100 μl DMEM with 10% FBS, were placed into the wells. After 4 h incubation, the numbers of migrated CD14<sup>+</sup> monocytes in the medium in the lower chamber were counted via flow cytometry. The numbers of migrated cells are expressed as cells per minute.

#### Macrophage polarization assay

Macrophage polarization assays were performed using Polycarbonate Membrane Transwell® Inserts (0.4-μm pores; Corning). CD14<sup>+</sup> monocytes ( $2 \times 10^5$ ) suspended in 600 µl RPMI-1640 (GIBCO) with 10% FBS were placed in the lower chambers. MSCs  $(2 \times 10^4)$  suspended in 100  $\mu$ l RPMI-1640 with 10% FBS were placed in the upper chambers. After 5 days incubation, flow cytometry was performed to determine the ratios of pro-inflammatory classically activated macrophages (M1 macrophages) or anti-inflammatory alternatively activated macrophages (M2 macrophages) in the lower chambers.

#### Enzyme-linked immunosorbent assay

MCP1 and TNF- $\alpha$  in cell culture supernatants or serum were measured using human MCP1 or TNF- $\alpha$  enzyme-linked immunosorbent assay (ELISA) kits according to the instruction manuals (BD OptEIA™).

### Alizarin red S (ARS) assay

MSCs were fixed and stained with 1% ARS for 15 min. Stained cells were observed via microscopy (40×). For ARS quantification, stained cells were incubated with 10% cetylpyridinium chloride monohydrate, and absorbance was measured at 562 nm.

#### Alkaline phosphatase assay

MSCs were fixed and treated with alkaline dye solution for 15 min in the dark. Stained cells were observed via microscopy (40×). To detect alkaline phosphatase (ALP) activity using ALP activity kits (Nanjing Jiancheng Biotech), MSCs were lysed using RIPA buffer containing protease and phosphatase inhibitors (Roche). Lysates were centrifuged and supernatants were incubated with reaction buffer at 37 °C for 15 min. After adding stop solution, absorbance was measured at 405 nm. Protein concentrations were measured using the BCA assay kit (Sigma-Aldrich). ALP activity is shown as units per gram protein per 15 min (U/gpro/15 min).

#### Lentiviruses assay

Four lentiviruses encoding short hairpin RNA (shRNA) for MCP1 (Lv-MCP1) and a negative control shRNA (Lv-NC) were constructed. Lentiviruses were generated via co-transfecting pGLVH1/GFP/Puro (GenePharma) and packing plasmids (pGag/Pol, pRev, and pVSV-G) into 293T cells. Cell culture supernatants containing lentiviruses were filtered and concentrated 72 h after transfection. Lentiviruses ( $10^9$  TU/ml) with 5  $\mu$ g/ml polybrene were incubated with MSCs for 24 h at a MOI of 50. Transduction efficiency was confirmed by green fluorescent protein (GFP)-positive MSCs and MCP1 ELISA. The most effective Lv-MCP1 with a target sequence of 5′-GGCT CGCGAGCTATAGAAGAA-3′ was chosen for experiments. The sequence for Lv-NC was 5′-TTCTCCGAACGTGT CACGTTTC-3′. Related experiments were performed on day 10 of induction.

#### ERK1/2 pathway blocking

U0126 (Sigma) was added (10 μM) as MSCs underwent osteogenic differentiation. Related experiments were performed on day 10 of induction.

#### Western blotting

Equal concentrations of proteins were separated via sodium dodecylsulfate-polyacrylamide gel electrophoresis and transferred to polyvinylidene fluoride membranes (Millipore), followed by blotting with primary antibodies against GAPDH, MCP1, C-C chemokine receptor 2 (CCR2), ERK1/2, and pERK1/2 (1:1000, Cell Signaling Technology). The membrane was washed and incubated with horseradish peroxidase (HRP)-conjugated secondary antibodies (1:3000, Cell Signaling Technology). Specific antibody-antigen complexes were detected using Immobilon Western Chemiluminescent HRP Substrate (Millipore).

#### Bone formation assay

In vivo bone formation assays were performed as described [\[26\]](#page-11-0). Briefly, MSCs  $(5 \times 10^5)$  were loaded onto 40 mg hydroxyapatite/tricalcium phosphate (HA/TCP; Zimmer) and implanted subcutaneously into the dorsal sides of 8 week-old BALB/c-nu/nu female mice (Laboratory Animal Center of Sun Yat-Sen University). Operations were performed under anesthesia achieved via intraperitoneal injection of ketamine and xylazine. At 2, 4, and 8 weeks, mice were killed and implants were obtained.

## Enthesial biopsies assay

Six AS patients and six non-AS patients (two patients with lumbar intervertebral disc herniation and four patients with scoliosis) were recruited. The sites of ossifying enthesis were confirmed via pre-surgery image analyses and visual observations during surgery. Ossifying tissues were obtained during lumbar spine surgeries. The characteristics of study subjects are presented in Supplemental Table 3.

## Immunofluorescence assay

Tissues obtained above were successively fixed, decalcified, and embedded in paraffin. Sections were deparaffinized, hydrated, and incubated in 1% Triton X-100/PBS. After antigen retrieval in citrate buffer and blocking in goat serum, sections were incubated with anti-MCP1 and anti-osteocalcin (OCN) antibodies (Abcam) overnight at 4 °C. Sections were washed and incubated with fluorescein secondary antibody and 4',6-diamidino-2phenylindole (DAPI; Thermo). Sections were examined with the Axio Observer fluorescence microscope (Carl Zeiss).

## Statistical analysis

All the results were determined based on at least three separate experiments containing at least triplicate samples. All data are expressed as means  $\pm$  standard deviations. T tests and one-way analyses of variance followed by Bonferroni tests and Pearson correlation tests were performed in SPSS (SPSS Inc). The n values indicated numbers of individuals in each experiment. P values less than 0.05 were considered statistically significant.

# **Results**

# ASMSCs secreted more MCP1 than HDMSCs during osteogenic differentiation

High-throughput microfluidics fluorescence qRT-PCR results show that MSCs expressed 27 main chemokines, with MCP1 expression increasing gradually, reaching a peak on day 10 of

induction, and then decreasing. ASMSCs expressed more MCP1 than HDMSCs on days 10 and 14 of induction. No differences were observed among the remaining 26 chemokines (Fig. [1a](#page-4-0)). Ordinary qRT-PCR produced consistent results (Fig. [1b](#page-4-0)). MCP1 protein expression in ASMSCs was higher than that in HDMSCs on days 10 and 14 of induction (Fig. [1c](#page-4-0), d). Besides, MCP1 secretion was positively correlated with ALP activity, a marker of early osteogenesis in MSCs, on day 10 of induction (Supplemental Fig. 1).

## ASMSCs outperformed HDMSCs in recruiting monocytes during osteogenic differentiation

Monocytes migrate and differentiate into macrophages, which subsequently polarize into either M1 or M2 macrophages [\[27](#page-11-0)]. MCP1 functions during monocyte migration and polarization  $[16]$  $[16]$  $[16]$ . CD14<sup>+</sup> monocytes were isolated, and their purity was confirmed (Supplemental Fig. 2). Corresponding to MCP1 expression results, monocyte migration induced by MSCs increased during MSC osteogenic differentiation, peaked on day 10 of induction and then decreased. ASMSCs had greater capacities to induce monocyte migration than HDMSCs on days 10 and 14 of induction (Fig. [2](#page-5-0)a). Monocyte migration induced by MSC culture supernatant showed similar results (Fig. [2b](#page-5-0)). We respectively isolated monocytes from healthy donors (HDMs) and AS patients (ASMs). However, regardless of whether monocyte migration was induced by MSCs directly or by MSC culture supernatant, only the source of MSCs, rather than the source of monocytes, had effects on migration (Fig. [2c](#page-5-0)). Moreover, no differences were observed in MCP1 and CCR2 expressions between HDMs and ASMs (Supplemental Fig. 3).

# Osteogenically differentiated ASMSCs exhibit reduced capacities to inhibit M1 macrophage polarization

Regardless of osteogenic differentiation status, MSCs inhibit monocytes polarization into M1 macrophages, which were positive for CD68 and HLA-DR. This capacity reduced on day 10 of induction. ASMSCs exhibited lower capacities than HDMSCs to inhibit M1 macrophage polarization on day 10 of induction. Surprisingly, no significant differences were observed in the effects of MSCs on M2 macrophage polarization, which were positive for CD68 and CD206 (Fig. [3](#page-6-0)a). Similar to migration assay results, HDMs and ASMs showed no differences in M1 macrophage polarization when cultured with the same MSCs on day 10 of induction (Fig. [3](#page-6-0)b). TNF- $\alpha$ is a pro-inflammatory cytokine primarily secreted by M1 mac-rophages [[28\]](#page-11-0). Similarly, MSCs inhibited TNF- $\alpha$  secretion from M1 macrophages, and this inhibition was reduced following osteogenic differentiation. On day 10 of induction, ASMSCs exhibited lower capacities than HDSMCs to inhibit TNF- $\alpha$  secretion (Fig. [3](#page-6-0)c).

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Fig. 1 ASMSCs secreted more MCP1 than HDMSCs during osteogenic differentiation. a Gene expression of chemokines was detected via highthroughput qRT-PCR. The heat maps show  $\Delta \Delta CT$  values. Each *column* represents one sample, and each row represents one gene. MCP1/CCL2 expression was higher in ASMSCs (A) than HDMSCs (H) during osteogenic differentiation. b MCP1 expression was confirmed via ordinary qRT-PCR. Results show that ASMSCs had higher MCP1

# Lv-MCP1 ameliorated dysfunctions in monocyte migration and polarization caused by osteogenically differentiated ASMSCs

Transfection and inhibition efficiencies were assessed in HDMSCs and ASMSCs (Fig. [4](#page-7-0)a). ARS and ALP assay results confirmed that ASMSCs had greater osteogenic differentiation capacities than HDMSCs. Lv-MCP1 did not affect osteogenic differentiation in either HDMSCs or ASMSCs (Fig. [4b](#page-7-0)). Moreover, Lv-MCP1 significantly decreased monocyte migration induced by HDMSCs and ASMSCs to equal levels on day 10 of induction (Fig. [4c](#page-7-0)). Similarly, inhibiting MCP1 in HDMSCs and ASMSCs significantly decreased M1 macrophages ratios to equal levels in the co-culture system (Fig. [4](#page-7-0)d). TNF- $\alpha$  secretion from M1 macrophages showed consistent results (Fig. [4e](#page-7-0)).



expression levels than HDSMCs on days 10 and 14 of induction. c ELISA results show that ASMSCs secreted more MCP1 than HDMSCs on days 10 and 14 of induction. d MCP1 expression at protein level was detected via Western blotting, confirming that ASMSCs expressed more MCP1 than HDMSCs on days 10 and 14 of induction. Data are presented as means  $\pm$  SD of 12 samples per group. \* indicates  $P < 0.05$  between HDMSCs and ASMSCs

# ERK1/2 pathway overactivation during abnormal osteogenic differentiation of ASMSCs leads to MCP1 overexpression and subsequent monocyte dysfunction

We previously demonstrated that BMP2 and Noggin imbalance led to ERK1/2 pathway overactivation, resulting in abnormal ASMSC osteogenic differentiation [[13\]](#page-10-0). Furthermore, ERK1/2 pathway modulates MCP1 expression [[29](#page-11-0)]. We confirmed that ERK1/2 pathway was more active in ASMSCs than HDMSCs on day 10 of induction. Adding U0126, a specific ERK1/2 inhibitor, reduced ERK1/2 pathway activation in HDMSCs and ASMSCs to equal levels (Fig. [5a](#page-8-0)). Moreover, U0126 completely inhibited osteogenic differentiation in HDMSCs and ASMSCs (Fig. [5b](#page-8-0)). On day 10 of induction, MCP1 secretion from HDMSCs and ASMSCs

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Fig. 2 ASMSCs outperformed HDMSCs in recruiting monocytes during osteogenic differentiation. Numbers of migrated CD14<sup>+</sup> monocytes were counted using flow cytometry. a MSCs promoted monocyte migration. ASMSCs  $(n = 12)$  showed a greater ability to promote monocyte migration than HDMSCs  $(n = 12)$  on days 10 and 14 of induction. **b** MSC culture supernatant promoted monocyte migration. ASMSCs  $(n = 12)$  culture supernatant exhibited a greater ability to promote

was significantly inhibited by U0126 and was consistent with that from undifferentiated MSCs (Fig. [5](#page-8-0)c). Blocking ERK1/2 pathway in HDMSCs and ASMSCs on day 10 of induction inhibited monocyte migration and decreased M1 macrophage ratios in the co-culture systems to equal levels (Fig. [5](#page-8-0)d, e). TNF- $\alpha$  secretion showed results consistent with macrophage polarization assays (Fig. [5](#page-8-0)f).

## MCP1 expression in MSCs during osteogenic differentiation in vivo

To detect MCP1 expression in MSCs during osteogenic differentiation in vivo, we constructed a mouse model bone formation assay [[26](#page-11-0)]. OCN expression in MSCs, a marker of osteogenesis, increased gradually from 2 to 8 weeks, indicating osteogenic differentiation. MCP1 expression in MSCs peaked at 4 weeks and decreased by week 8. Both the expression of MCP1 in ASMSCs were markedly higher than were observed for HDMSCs at 4 weeks (Fig. [6a](#page-9-0), white arrows).

monocyte migration than HDMSCs  $(n = 12)$  culture supernatant on days 10 and 14 of induction. c Monocytes from healthy donors (HDMs;  $n = 12$ ) and monocytes from AS patients (ASMs;  $n = 12$ ) showed equal migration capacities when induced by the same types of MSCs or culture supernatants on day 10 of induction. The experiments were done in triplicate and data are presented as means  $\pm$  SD. \* indicates  $P < 0.05$ between HDMSCs and ASMSCs

## MCP1 expression at local sites of ossifying enthesis in AS patients

MCP1 expression at local sites of ossifying enthesis in vivo in AS has never been investigated. Therefore, we evaluated MCP1 expression in osteoblasts, which come from MSCs and are major cells responsible for new bone formation [[30\]](#page-11-0), at local sites of ossifying enthesis in AS. The expression of  $MCP1$  in  $OCN<sup>+</sup>$  osteoblasts was significantly higher in AS patients than in non-AS patients (Fig. [6](#page-9-0)b, white arrow heads). Serum MCP1 concentrations in AS patients were essentially the same as in healthy donors (Supplemental Fig. 4).

#### Discussion

MCP1 is a crucial chemokine for maintaining immunity homeostasis [\[16](#page-10-0)]. Previous studies demonstrated that serum MCP1 concentrations are elevated in patients with inflammatory diseases [\[31](#page-11-0), [32](#page-11-0)]. However, whether MCP1 levels are elevated in AS is controversial [[18](#page-10-0), [33](#page-11-0)]. Here, we

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Fig. 3 Osteogenically differentiated ASMSCs exhibit reduced capacities to inhibit M1 macrophage polarization. a M1 macrophage polarization ratios decreased after culturing with MSCs. M1 macrophage polarization ratios in cultures with ASMSCs  $(n = 12)$  were equal to those with HDMSCs  $(n = 12)$  on day 0 of induction. M1 macrophage polarization ratio with ASMSCs  $(n = 12)$  was higher than those with HDMSCs  $(n = 12)$  on day 10 of induction. M2 macrophage polarization ratios were almost not affected by HDMSCs  $(n = 12)$  or ASMSCs  $(n = 12)$ . **b** When cultured with the same MSCs on day 10 of induction, M1

macrophage polarization ratios of HDMs ( $n = 12$ ) showed no statistical differences with those of ASMs ( $n = 12$ ). c TNF- $\alpha$  secretion by macrophage decreased after culturing with MSCs. TNF- $\alpha$  secretion of macrophage cultured with ASMSCs  $(n = 12)$  was equal to that of those cultured with HDMSCs ( $n = 12$ ) on day 0 of induction. TNF- $\alpha$  secretion of macrophage cultured with ASMSCs  $(n = 12)$  was higher than those cultured with HDMSCs  $(n = 12)$  on day 10 of induction. The experiments were done in triplicate and data are presented as means ± SD. \* indicates  $P < 0.05$  between HDMSCs and ASMSCs

demonstrated that serum MCP1 levels are equal between healthy donors and AS patients. However, MCP1 expression in osteoblasts at sites of ossifying enthesis in vivo was higher in AS patients than in non-AS patients, indicating that MCP1 is locally, rather than systemically, elevated in AS. MSCs are primary sources of osteoblasts and one of the main producers

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Fig. 4 Lv-MCP1 ameliorated dysfunctions in monocyte migration and polarization caused by osteogenically differentiated ASMSCs. a Transduction efficiency in MSCs was confirmed based on GFP-positive cells and ELISA results. Scale  $bar = 200 \mu m$ . b ARS and ALP assay results show that ASMSCs  $(n = 9)$  had greater osteogenic differentiation capacity than HDSMCs  $(n = 9)$  on day 10 of induction. Lv-MCP1 did not alter osteogenic differentiation capacity of MSCs. Scale bar = 50  $\mu$ m. c The capacities of HDMSCs  $(n = 9)$  and ASMSCs  $(n = 9)$  to induce monocyte migration on day 10 of induction were decreased to equal

levels by Lv-MCP1. **d** The capacities of HDMSCs  $(n = 9)$  and ASMSCs  $(n = 9)$  to inhibit M1 macrophage polarization on day 10 of induction were increased to equal levels by Lv-MCP1. e The capacities of HDMSCs  $(n = 9)$  and ASMSCs  $(n = 9)$  to inhibit TNF- $\alpha$  secretion on day 10 of induction were increased to equal levels by Lv-MCP1. The experiments were done in triplicate and data are presented as means  $\pm$  SD. \* indicates  $P < 0.05$  between HDMSCs and ASMSCs. & indicates  $P < 0.05$  for Lv-MCP1 group compared with induction group. Control indicates MSCs cultured in growth medium

of MCP1 [[30](#page-11-0), [34](#page-11-0)]. Besides, MSCs continuously secrete MCP1 during osteogenic differentiation [\[23\]](#page-10-0). Therefore, we hypothesized that MCP1 overexpression in osteoblasts at local osteogenesis sites in AS patients derives from ASMSCs undergoing abnormal osteogenic differentiation. Supporting this hypothesis, we showed that ASMSCs secreted more MCP1 than HDMSCs during osteogenic differentiation both in vitro (10 days) and in vivo (4 weeks) at time points corresponding to early stages of differentiation. It is acknowledged that MCP1 parasecretion from MSCs results in immune dysfunction [\[22](#page-10-0)]. Whether MCP1 overexpression during abnormal osteogenic differentiation of ASMSCs leads to subsequent immunocyte dysfunction remains unknown.

MCP1 has various effects on monocytes, one of which is inducing monocyte migration [\[16\]](#page-10-0). Under physiological conditions, MSCs recruit monocytes for tissue repair by secreting MCP1 [\[8\]](#page-10-0). However, MCP1 overexpression in MSCs accelerates monocyte migration and results in pathological conditions [\[35\]](#page-11-0). We observed that the capacity of ASMSCs to recruit monocytes increased during osteogenic differentiation and corresponded to MCP1 overexpression. Moreover, Lv-MCP1 could ameliorate this dysfunction. These results suggest that the enhanced ability of ASMSCs to induce monocyte migration results from MCP1 overexpression during abnormal osteogenic differentiation.

MCP1 affects monocyte polarization and cytokine secretion after migration [\[16](#page-10-0), [36](#page-11-0)]. Although it has been demonstrated that MSCs inhibit M1 macrophage polarization [\[37\]](#page-11-0), we found that this capacity was reduced after osteogenic differentiation. Moreover, osteogenically differentiated ASMSCs had lower capacities to inhibit M1 macrophage polarization than HDMSCs. This dysfunction, along with aberrant induction of monocyte migration, was not only consistent with MCP1 overexpression patterns but also ameliorated by Lv-MCP1. These results suggest that the above changes resulted from MCP1 overexpression. Actually, the effects of MCP1 on monocyte polarization remain controversial. Some studies concluded that MCP1 promotes monocyte polarization into M2 macrophages rather than M1 macrophages [[36](#page-11-0), [38](#page-11-0)]. Conversely, other studies, consistent with our results,

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Fig. 5 ERK1/2 pathway overactivation during abnormal osteogenic differentiation of ASMSCs leads to MCP1 overexpression and subsequent monocyte dysfunction. a Western blotting showed that ASMSCs  $(n = 9)$  had higher ERK1/2 levels than HDMSCs  $(n = 9)$  on day 10 of induction. U0126 decreased ERK1/2 pathway activation in HDMSCs  $(n = 9)$  and ASMSCs  $(n = 9)$  to equal levels. **b** ARS and ALP assays confirmed the greater osteogenic differentiation capacity of ASMSCs  $(n = 9)$  relative to HDMSCs  $(n = 9)$  on day 10 of induction, which was decreased to equal levels by U0126. Scale bar = 50  $\mu$ m. c MCP1 secretion levels were higher in ASMSCs  $(n = 9)$  than in HDMSCs  $(n = 9)$  on day 10 of induction and were decreased to equal levels following treatment with U0126. **d** ASMSCs  $(n = 9)$  had a greater

indicated that MCP1 is positively related to M1 macrophage ratio and promotes pro-inflammatory cytokine secretion [\[39](#page-11-0)–[41\]](#page-11-0). These discrepancies may result from differences in the microenvironments, culture systems, and cell origins used in each study. TNF- $\alpha$ , mainly secreted by M1 macrophages, is the most important inflammatory cytokine in AS [[42](#page-11-0)]. In this study, TNF- $\alpha$  secretion patterns were fully consistent with M1 macrophage polarization, further supporting to our conclusions and providing insights into why high levels of TNF- $\alpha$ exist at specific locations in AS.

MCP1 can be secreted by monocytes and exerts effects by binding to CCR2, a specific receptor for MCP1 [[16,](#page-10-0) [36](#page-11-0)]. However, no differences in MCP1 and CCR2 expression were observed between HDMs and ASMs. Besides, HDMs and ASMs exhibited similar migration and polarization capabilities when cultured with the same MSCs. Moreover, inhibiting

capacity to induce monocyte migration than HDMSCs  $(n = 9)$  on day 10 of induction, but these capacities were equalized by U0126. e ASMSCs  $(n = 9)$  had a lower capacity to inhibit M1 macrophage polarization than HDMSCs  $(n = 9)$  on day 10 of induction. This was restored by treatment with U0126. f The capacity to inhibit TNF- $\alpha$ secretion was lower in ASMSCs  $(n = 9)$  than in HDMSCs  $(n = 9)$  on day 10 of induction but was equalized to equal levels by U0126. The experiments were done in triplicate and data are presented as means ± SD.  $*$  indicates  $P < 0.05$  between HDMSCs and ASMSCs.  $\&$  indicates  $P < 0.05$  for OM group compared with OM with U0126 group. OM indicates osteogenic differentiation medium

MCP1 expression in HDMSCs and ASMSCs removed any differences in monocyte migration and polarization. These results suggest that MCP1 overexpression during abnormal osteogenic differentiation of ASMSCs triggers monocyte dysfunctions.

ERK1/2 pathway contributes to osteogenesis [[43](#page-11-0)]. Particularly, its activation promotes MSC osteogenic differentiation [\[44](#page-11-0)]. We previously demonstrated that an imbalance between BMP2 and Noggin resulted in excessive ERK1/2 pathway activation, leading to abnormal osteogenic differentiation of ASMSCs. Decreasing BMP2 or increasing Noggin rectified ERK1/2 pathway overactivation and abnormal osteogenic differentiation of ASMSCs [\[13](#page-10-0)]. Additionally, ERK1/2 pathway activation leads to MCP1 secretion [\[29](#page-11-0), [45\]](#page-11-0). We found that blocking ERK1/2 pathway inhibited abnormal osteogenic differentiation of ASMSCs and reduced the

<span id="page-9-0"></span>Fig. 6 MCP1 expression in MSCs during osteogenic differentiation in vivo and MCP1 expression at local sites of ossifying enthesis in AS patients. a OCN expression increased gradually from 2 to 8 weeks, indicating the osteogenic differentiation of MSCs. MCP1 expression of MSCs in the scaffolds peaked at 4 weeks and then decreased at 8 weeks. The expressions of MCP1 in ASMSCs were markedly higher than were observed for HDMSCs at 4 weeks (white arrows). b MCP1 expression was higher in OCN+ osteoblasts at sites of ossifying enthesis in AS patients compared with non-AS patients (white arrow heads). S indicates scaffold; M indicates MSCs. Scale bar =  $50 \mu m$ 



subsequent MCP1 overexpression, indicating that MCP1 overexpression results from ERK1/2 pathway overactivation during abnormal osteogenic differentiation. These conclusions not only emphasize the key role of ERK1/2 pathway in both abnormal osteogenic differentiation and MCP1 overexpression but also illustrate an inherent relationship between these two abnormalities in ASMSCs. Evidentially, a positive correlation was found between ALP activity and MCP1 expression in MSCs. However, Lv-MCP1 did not alter osteogenic differentiation capacity in HDMSCs or ASMSCs, indicating that MCP1 overexpression does not cause abnormal osteogenic differentiation of ASMSCs. We speculate that MCP1 overexpression results from abnormally osteogenic differentiation of ASMSCs. Therefore, we further conclude that the effects of MCP1 overexpression on monocyte migration and polarization are in order the result of abnormal ASMSC osteogenic differentiation.

The relationship between pathological osteogenesis and inflammation in AS remain controversial [\[2\]](#page-10-0). Previous studies found that CD14<sup>+</sup> monocytes and CD68<sup>+</sup> macrophages infiltrate close to the area of new bone formation in AS [\[42,](#page-11-0) [46\]](#page-11-0). This in vivo phenomenon is similar to the in vitro results of our study. Do inflammatory cells infiltrate before or after osteogenesis? To the best of our knowledge, whether pathological osteogenesis leads to inflammation remains unknown in AS. We previously demonstrated that ASMSCs have enhanced capacities for osteogenic differentiation [[13](#page-10-0)]. We further demonstrated that ASMSCs secreted more MCP1 during abnormal osteogenic differentiation, resulting in monocyte dysfuncions that are the typical of inflammation in AS [\[42\]](#page-11-0). Therefore, we propose a novel mechanism that pathological osteogenesis can lead to inflammation in AS. During abnormal osteogenic differentiation, MCP1 secretion by ASMSCs greatly increases, which recruits additional monocytes from circulating blood to osteogenesis sites. These <span id="page-10-0"></span>recruited monocytes polarize into M1 macrophages and secrete TNF- $\alpha$  under the effects of MCP1, leading to local inflammation that is observed in AS (Supplemental Fig. 5). This mechanism may suggests a vicious cycle between pathological osteogenesis and inflammation, and pathological osteogenesis may be one of the dynamic factors of inflammation in AS. Furthermore, this mechanism explains why inhibiting inflammation alone does not delay pathological osteogenesis [7], and it indicates that simultaneous treatments targeting pathological osteogenesis and inflammation may have surprising benefits for AS patients.

Here, we demonstrated that MCP1 triggers inflammation during abnormal osteogenic differentiation of MSCs in AS, providing insights into AS pathogenesis and the relationship between pathological osteogenesis and inflammation. However, several limitations exist in this study. How does the complex in vivo environment affect MSC osteogenic differentiation and monocyte function? Continuous imaging studies and histological examinations are required to support this hypothesis, which should be addressed in future studies.

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Compliance with ethical standards This study conforms to the Declaration of Helsinki and was approved by the Ethics Committee and the Animal Ethical and Welfare Committee of Sun Yat-Sen Memorial Hospital, Sun Yat-Sen University, Guangzhou, China.

Conflict of interest The authors declare that they have no conflicts of interests.

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