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# The secretome of equine bone marrow-derived mesenchymal stem cells enhanced regenerative phenotype of equine articular chondrocytes





Manon Jammes<sup>1,†</sup>, Romain Contentin<sup>1,†</sup>, Frédéric Cassé<sup>1</sup>, Bastien Bourdon<sup>1</sup>, Arnaud Bianchi<sup>2</sup>, Fabrice Audigié<sup>3</sup>, Thomas Branly<sup>1</sup>, Emilie Velot<sup>2</sup>, and Philippe Galéra<sup>1</sup>

<sup>1</sup> Normandie Univ, UNICAEN, BIOTARGEN, F-14000 Caen, France.

<sup>2</sup> University of Lorraine, CNRS, IMoPA, F-54000 Nancy, France.

<sup>3</sup> Center of Imaging and Research on Locomotor Affections on Equines (CIRALE), Unit Under Contract 957 Equine Biomechanics and Locomotor Disorders (USC 957 BPLC), French National Research Institute for Agriculture Food and Environment (INRAE), Ecole Nationale Vétérinaire d'Alfort, F-94700 Maisons-Alfort, France.

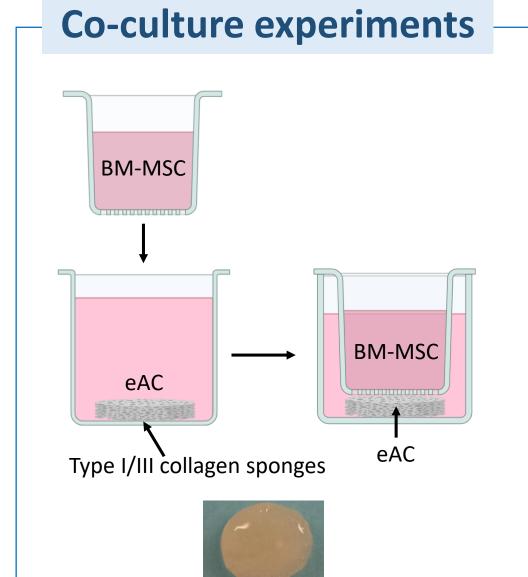
† These authors contributed equally to this work.

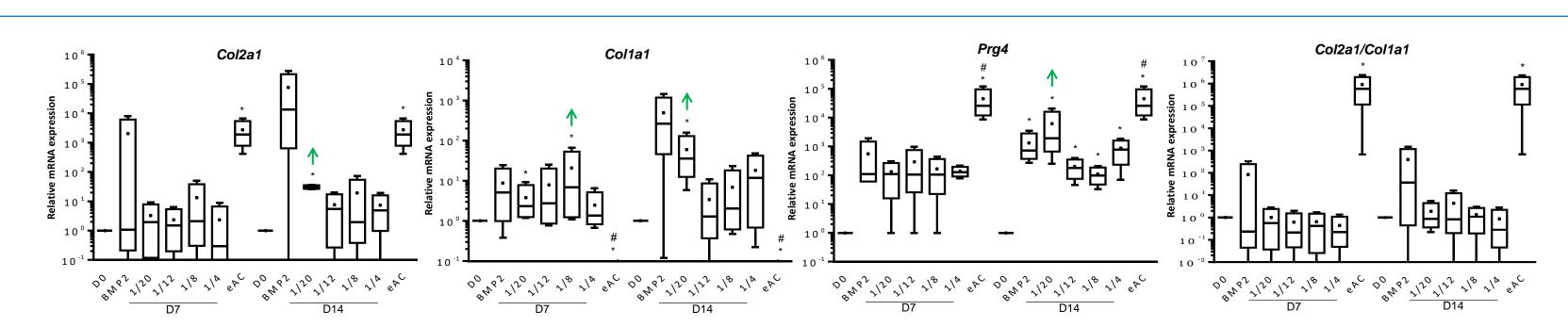
## Introduction

Equine **osteoarthritis** (OA) is a sequential disease which leads to cartilage degradation and painful bone frictions. It induces impaired animal well-being, premature cessation of sport activity, and financial losses. Fibrocartilage synthesis occurring during cartilage destruction is a physiological response, allowing bone protection but reducing tissue mechanical resistance. To date, there is a **lack of curative therapies**. Mesenchymal stem cell (MSC)-based therapies are promising for cartilage repair, but face limitations inherent to cell itself which can be overcome using their **secretome** through acellular therapy approaches.

To understand the effects of equine bone marrow (BM-)MSC secretome on equine articular chondrocytes (eAC) phenotype, indirect co-culture experiments were first performed. Then, we wanted to recapitulate the effects we observed in co-culture on eAC using the MSC-conditioned medium (CM) to make acellular therapy conceivable. We assessed hyaline cartilage and fibrocartilage markers at the transcription and protein levels, and evaluated eAC migratory capacities, which are of interest during OA therapy to favor the filling of the cartilage defects. To optimize immunomodulation properties of MSC secretome for future experiments, MSC priming relevance with interleukin (IL)1-β was evaluated. Suspected to be the principal vectors of the effects observed, exosomes were isolated through chemical precipitation and then characterized to confirm their nature.

# Results





Cartilage markers

Col2a1: Hyaline cartilage

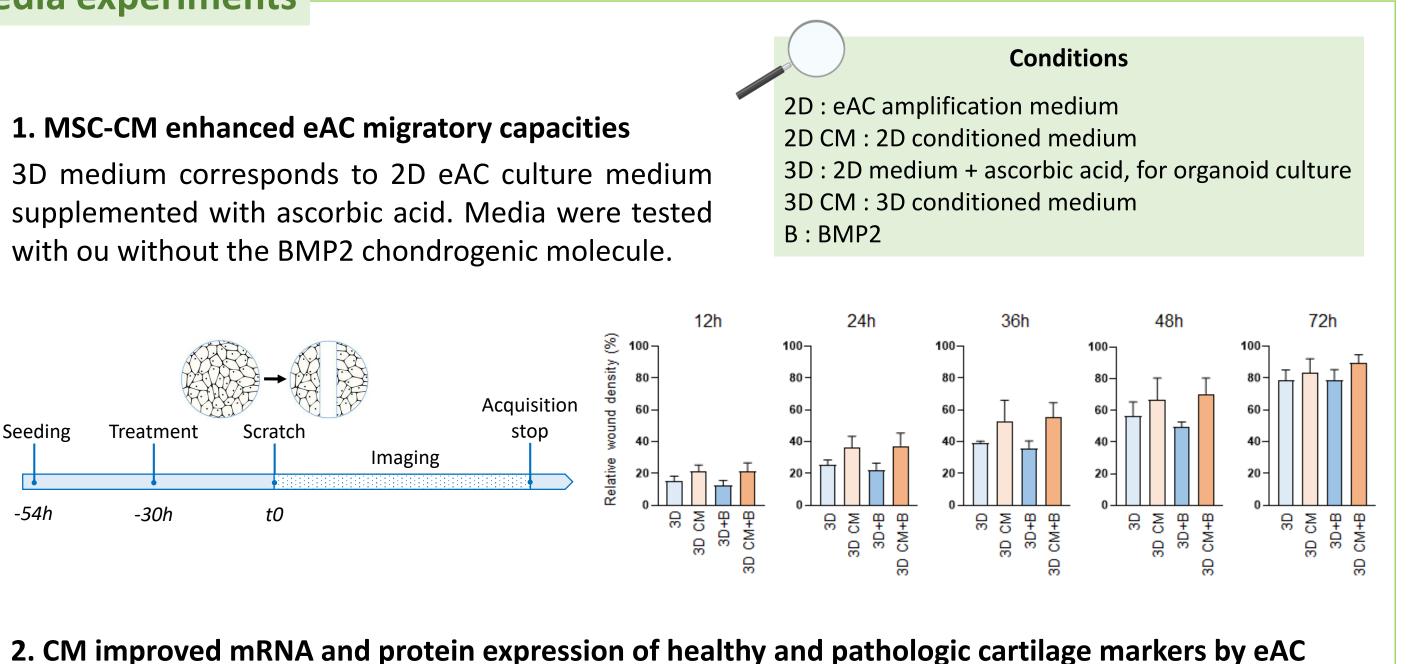
Col1a1: Fibrocartilage, less resistant than type II collagen

Prg4: Lubricin, joint lubricant

mRNA levels of hyaline cartilage and fibrocartilage matrix markers were assessed by RT-qPCR after 7 (D7) and 14 days (D14) of culture. Normalization was performed according to D0 (eAC cultured in monolayer until P3). The significance of the values was tested using a Mann-Withney test (\* p<0,05). A chondrogenic control was performed using bone morphogenetic protein 2 (BMP2), and eAC condition refers to mRNA extracted from eAC at P0.

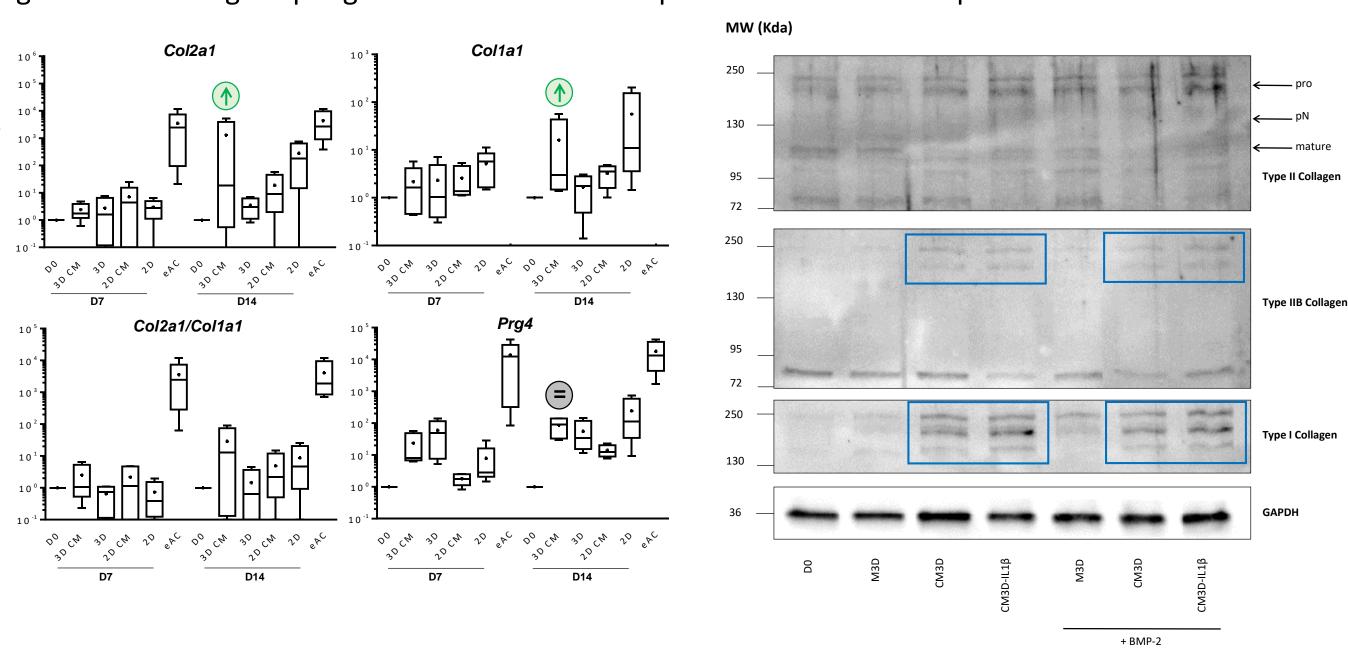
eAC co-culture with BM-MSC improved eAC phenotype by enhancing *Col2a1*, *Col1a1*, *Prg4* markers expressions, but *Col2a1/Col1a1* remained unchanged. 
→ BM-MSC secretome modulates eAC phenotype

# Conditioned media experiments 1. MSC-CM enhanced 3D medium corresponsupplemented with as with ou without the BN Seeding Treatment Scratch (0.22 μm) Aliquoting Conditioned media (CM) collection Seeding Treatment Scratch (0.22 μm) After 7 and 14 days of and fibrocartilage macondition. CM3D-IL1β condition prior to conditioning organoid-like collagen sorganoid-like sorganoid-like collagen sorganoid-like collagen sorganoid-like co



2. CM improved mRNA and protein expression of healthy and pathologic cartilage markers by eAC After 7 and 14 days of 2D and 3D culture of eAC with MSC-CM, mRNA expression of hyaline cartilage and fibrocartilage markers was assessed by RT-qPCR. Normalization was realized towards the D0 condition.

CM3D-IL1β condition refers to CM from MSC primed with interleukine(IL)-1β (10 ng/mL) during 24h prior to conditioning. After 14 days, eAC cultured with CM supplemented or not with BMP2, in organoïd-like collagen sponges were harvested and protein extraction was performed.



MSC-CM enhanced hyaline cartilage and fibrocartilage markers expression of eAC after 14 days and increased their migratory capacities

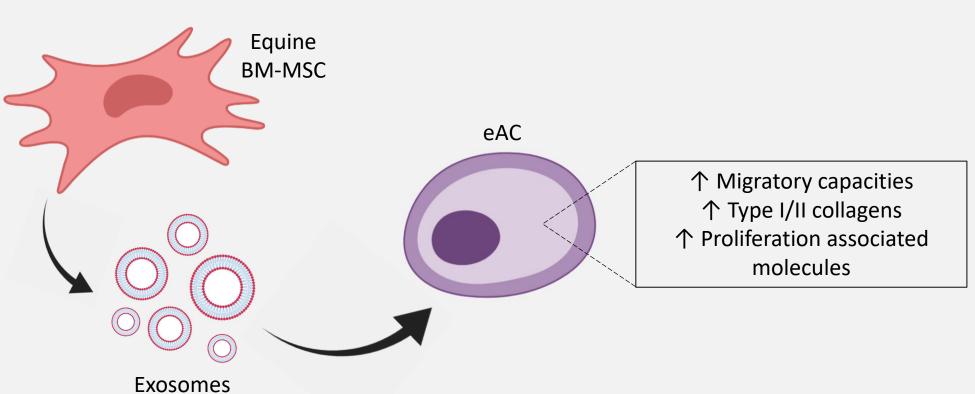
# → BM-MSC-CM recapitulates the effect of co-culture on eAC

MSC secretome was able to increase collagen protein accumulation especially when primed with the **OA-related inflammatory cytokine IL-1\beta**, suggesting that the therapeutic potential of MSC-CM could be enhanced by an optimized cytokine priming.

### **Exosomes characterization** 1. Physical and morphological 2. Expression of exosomal characterization specific markers MSC-CM nanoparticle size extraction Protein distribution and concentration performed after purification were determined by NTA (A). of exosomes according to Particle morphology the detailed protocol. Then, observed by TEM to confirm western-blot their EV nature and assess EV performed with antibodies integrity (B). exosome-specific against proteins. Elimination of membrane Average size debris and large particles 165.1 ± 1.8 nm $4.19x10^9 \pm 1.64x10^8$ particles/mL Filtration (0.22 μm) Concentration by Amicon filtration (cut-off: 100 kDa) PBS washing **Exosomes precipitation** PEG 50% Protamine sulfate 2% 3. Uptake of exosomes by eAC GAPDH CM harvest Agitation 1h 24h incubation Concentration Fluorescence imaging 12 000g 10 mn CM from BM-MSC contains undamaged exosomes RIPA-based expressing Alix/TSG101/CD81. These extracellular lysis for protein extraction vesicles interact with eAC and can be uptaken by the cells.

# Conclusions

eAC 3D culture



MSC-CM were able to favor eAC migratory capacities and cartilage matrix synthesis *in vitro* by increasing types II and I collagens protein amounts. These effects could be mediated by exosomes since we have demonstrated that MSC-CM contain these nanovesicles that can be uptaken by eAC. However further investigations are needed to better define the contribution of exosomes to the therapeutic potential of MSC-CM in OA.

These results demonstrate that **MSC secretome influences eAC phenotype** and could relieve OA symptoms or delay OA outcomes in equine medicine. Consequently, an acellular therapy avoiding disadvantages of direct cell use must be considered, especially because this process could be potentially transposed to human medicine according to the One Health concept.

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