# Interaction of HIF1 $\alpha$ and $\beta$ -catenin inhibits matrix metalloproteinase 13 expression and prevents cartilage damage in mice

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Low oxygen tension (hypoxia) regulates chondrocyte differentiation and metabolism. Hypoxia-inducible factor  $1\alpha$  (HIF1 $\alpha$ ) is a crucial hypoxic factor for chondrocyte growth and survival during development. The major metalloproteinase matrix metalloproteinase 13 (MMP13) is also associated with chondrocyte hypertrophy in adult articular cartilage, the lack of which protects from cartilage degradation and osteoarthritis (OA) in mice. MMP13 is up-regulated by the Wnt/β-catenin signaling, a pathway involved in chondrocyte catabolism and OA. We studied the role of HIF1 $\alpha$  in regulating Wnt signaling in cartilage and OA. We used mice with conditional knockout of  $Hif1\alpha$ ( $\Delta$ Hif1 $\alpha$ <sup>chon</sup>) with joint instability. Specific loss of HIF1 $\alpha$  exacerbated MMP13 expression and cartilage destruction. Analysis of Wnt signaling in hypoxic chondrocytes showed that HIF1 $\alpha$  lowered transcription factor 4 (TCF4)- $\beta$ -catenin transcriptional activity and inhibited MMP13 expression. Indeed, HIF1 $\alpha$  interacting with  $\beta$ -catenin displaced TCF4 from *MMP13* regulatory sequences. Finally,  $\Delta Hif1\alpha^{chon}$  mice with OA that were injected intraarticularly with PKF118-310, an inhibitor of TCF4-\beta-catenin interaction, showed less cartilage degradation and reduced MMP13 expression in cartilage. Therefore, HIF1 $\alpha$ - $\beta$ -catenin interaction is a negative regulator of Wnt signaling and MMP13 transcription, thus reducing catabolism in OA. Our study contributes to the understanding of the role of HIF1 $\alpha$  in OA and highlights the HIF1 $\alpha$ - $\beta$ -catenin interaction, thus providing new insights into the impact of hypoxia in articular cartilage.

hypoxia-inducible factor  $1\alpha$  | chondrocyte | osteoarthritis | Wnt signaling | matrix metalloprotease 13

ow oxygen tension (hypoxia) orchestrates several cell functions and is critical in health and disease (1–4). Hypoxiainducible factor  $1\alpha$  (HIF1 $\alpha$ ) is an essential factor to maintain chondrocyte homeostasis and allow cell differentiation (5, 6). HIF1 is a heterodimeric DNA-binding complex containing a constitutive HIF1 $\beta$  subunit and HIF1 $\alpha$  subunit. In hypoxia, HIF1 binds to the hypoxia response elements of target genes, whereas in normoxia, HIF1 $\alpha$  is hydroxylated, thereby leading to its degradation. Indeed, HIF1 $\alpha$  hydroxylation is recognized by the von Hippel–Lindau tumor suppressor protein (pVHL), an E3 ubiquitin ligase that targets HIF $\alpha$  for proteolysis in the proteasome (7). The HIF1 $\alpha$  pathway interacts with different cell signaling pathways, among them Wnt signaling. Indeed, HIF1a interacts with β-catenin in regulating cell growth and survival. In embryonic stem cells, HIF1 $\alpha$ - $\beta$ -catenin complexes up-regulate lymphoid enhancer-binding factor 1 and transcription factor 1 (TCF1), which activates Wnt signaling (8), whereas in colorectal cancer cells, HIF1 $\alpha$  blocks the TCF4- $\beta$ -catenin interaction and transcriptional activity, thus inhibiting canonical Wnt signaling (9).

Cartilage loss characterizes osteoarthritis (OA), one of the most frequent joint disorders, but available treatments are poorly efficient to prevent joint destruction (10, 11). Therefore, the need for novel drug targets to treat OA is paramount. Matrix metalloproteinase 13 (MMP13) triggers the degradation of articular cartilage. Indeed, chondrocyte-specific deletion of MMP13 alleviated OA in mice; the Wnt family members were candidates for the regulation of MMP13 expression in chondrocytes because its expression was increased in chondrocytes from mice with conditional activation of  $\beta$ -catenin (12). Cumulative data showed that Wnt activity is low under physiological conditions, and activation of Wnt signaling contributes to cartilage breakdown in OA (13, 14). The modulation of Wnt inhibitors had significant effects on chondrocyte catabolism of mice. Indeed, loss of sclerostin enhanced cartilage degradation (15) and the over-expression of Dkk-1 alleviated OA (14). Despite the hypoxic status of cartilage (16), the involvement of hypoxia in regulating Wnt signaling and MMP13 expression in cartilage is still unclear.

We studied the role of HIF1 $\alpha$  in regulating Wnt signaling in cartilage of mice with conditional knockout of *Hif1* $\alpha$  ( $\Delta$ *Hif1* $\alpha$ <sup>*chon*</sup>) and induced OA. Hypoxia maintained low Wnt/ $\beta$ -catenin signaling via HIF1 $\alpha$ , which lowered MMP13 expression, then prevented chondrocyte catabolism and cartilage loss. Here we therefore highlight the role of HIF1 $\alpha$  in cartilage remodeling and loss during OA and provide a previously unidentified mechanism of micro-environmental regulation of Wnt signaling in cartilage.

## Results

HIF1 $\alpha$  Deletion in Chondrocytes Enhanced OA Development and MMP13 Expression in Mice. Articular chondrocytes are physiologically in a hypoxic state that might be altered in OA (5). To

# Significance

Hypoxia-inducible factor  $1\alpha$  (HIF1 $\alpha$ ) is important for cell growth and survival. It modulates Wnt signaling, regulating cell differentiation and fate. Osteoarthritis (OA) is an increasingly frequent joint disorder characterized by progressive cartilage breakdown in which Wnt/ $\beta$ -catenin signaling triggers matrix metalloproteinase 13 (MMP13) expression and chondrocyte catabolism. Here we demonstrate HIF1 $\alpha$  inhibits  $\beta$ -catenin signaling by blocking transcription factor 4 (TCF4)– $\beta$ -catenin interaction and down-regulates MMP13 expression, thereby alleviating cartilage lesions, whereas the TCF4–  $\beta$ -catenin signaling induces an OA phenotype in mice. In OA joints, PKF118-310, a small molecule that blocked TCF4– $\beta$ -catenin interaction, significantly reduced the progression of OA cartilage lesions. Thus, blockade of TCF4– $\beta$ -catenin signaling by HIF1 $\alpha$  represents a promising strategy to prevent articular cartilage loss in OA.

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monitor oxygen tensions in OA chondrocytes, we administered a hypoxyprobe (pimonidazole hydrochloride) to control and OA mice. Hypoxia levels were markedly decreased in all layers of the OA articular cartilage, as shown by immunohistochemistry and by the decreased number of hypoxic cells (Fig. 1.4). HIF1 $\alpha$  was expressed in undamaged articular cartilage at baseline but its level decreased significantly in OA joints along with the increased cartilage damage (Fig. 1*B*). Noteworthy, the number of Hif1 $\alpha$ -expressing cells was steadily down-regulated after destabilization of the medial meniscus (4 and 6 wk), whereas OA damage increased in a time-dependent manner.

To determine the role of HIF1 $\alpha$  in OA development, we generated mice with inducible conditional knockout of  $Hif1\alpha$  by mating Col2- $Cre^{ERT}$  mice with  $Hif1\alpha^{Il/I}$  mice in which the recombination was induced by tamoxifen. We first verified that the *Cre-lox* recombination occurred correctly in cartilage in *Col2*- $Cre^{ERT}$ ; R26R-*LacZ* mice and R26R-*LacZ* mice were used as controls.  $\beta$ -Galactosidase was expressed in the articular cartilage of *Col2*- $Cre^{ERT}$ ; R26R-*LacZ* mice, thus tamoxifen induced *Cre-lox* recombination in chondrocytes (Fig. S1). OA was induced in  $\Delta Hif1\alpha^{chon}$  mice 1 wk after tamoxifen injections. OA cartilage

lesions were increased in  $\Delta Hif1\alpha^{chon}$  mice as shown by the osteoarthritis score (Fig. 1*C*). As HIF1 $\alpha$  is a survival factor, we assessed chondrocyte apoptosis in OA in the absence of HIF1 $\alpha$ . The number of apoptotic cells was increased in  $\Delta Hif1\alpha^{chon}$  mice although the OA score remained unchanged (Fig. 1*D*). TUNEL positive cells were further increased in  $\Delta Hif1\alpha^{chon}$  mice with destabilization of the medial meniscus (DMM). As previously described, the expression of MMP13 was induced in OA mice. This increase is enhanced in  $\Delta Hif1\alpha^{chon}$  mice along with the exacerbated cartilage loss (Fig. 1*E*). Thus, OA is associated with loss of hypoxia in articular cartilage, decreased HIF1 $\alpha$  protein levels, and increased MMP13 expression and cartilage loss in mice. To rule out the hypothesis that  $\Delta Hif1\alpha^{chon}$  could induce endothelial PAS domain-containing protein 1 (EPAS1 or Hif2 $\alpha$ ) expression and therefore contribute to the phenotype, we found that EPAS1 was expressed at the same level in  $Hif1\alpha^{fl0T}$  mice and  $\Delta Hif1\alpha^{chon}$  mice, suggesting the absence of compensatory increase of EPAS1 (Fig. S24).

HIF1 $\alpha$  Inhibits *MMP13* Expression and the Transcription of Wnt Targets. HIF1 $\alpha$  is a major HIF that regulates chondrocyte metabolism (17). We first assessed the role of hypoxia in the



Fig. 1. Hypoxia and hypoxia-inducible factor  $1\alpha$ (HIF1 $\alpha$ ) are reduced in cartilage of mice with OA. (A) Immunohistofluorescence staining of Hypoxyprobe adducts in healthy and OA wild-type mouse cartilage. Hypoxyprobe adducts were revealed in hypoxic cells ( $P_{O2}$  < 10 mmHg) by a fluorescein-conjugated antibody (HP-FITC-MAb). Graph shows proportion of hypoxic positive cells in cartilage of the tibial plateau and internal femoral condyle. Data are mean  $\pm$  SEM. \*P < 0.05 compared with control (n = 7 animals per group). (Scale bar, 100 μm.) (B, Upper) HIF1α immunostaining and OA score (Safranin-O staining) in control mice at 0, 4, and 6 wk post-OA induction. Graphs show percentage of HIF1 $\alpha$ (+) cells and OA score in articular cartilage of the tibial plateau and internal femoral condyle. (Scale bar, 100  $\mu$ m.) \*P < 0.05 compared with control (n = 7 animals per group) (C) Safranin-O staining and (D) TUNEL assay of  $Hif1\alpha^{flifl}$  and  $\Delta Hif1\alpha^{chon}$  mouse joints with OA or sham operation (ct) at week 6. (Scale bar, 100 µm.) OA score in OA and sham-operated knees of  $Hif1\alpha^{fl/f}$ and  $\Delta Hif1 a^{chon}$  mice (week 6). \*P < 0.05 compared with control.  ${}^{\#}P < 0.05$  (n = 8-11 animals per group). (E) Immunostaining for MMP13 in  $Hif1\alpha^{flfl}$  and  $\Delta Hif1 \alpha^{chon}$  mouse joints (week 6) and quantification. (Scale bar, 100  $\mu$ m.) \*P < 0.05 compared with control.  ${}^{\#}P < 0.05 \ (n = 8-11 \text{ animals per group}).$ 

metabolic effects of Wnt signaling in primary chondrocytes. As expected in normoxia, Wnt3a reduced proteoglycan release and *Col2A* expression while increasing *Mmp13* expression (Fig. 2 *A* and *B*). In contrast, under hypoxic conditions, Wnt3a failed to modulate the proteoglycan release as well as the expression of catabolic markers. Furthermore target genes such as *Axin* and *Wisp1* were not regulated by Wnt3a in hypoxia (Fig. 2C). To investigate whether HIF1 $\alpha$  modulates *MMP13* expression induced by Wnt, we first deleted HIF1 $\alpha$  chondrocytes and analyzed the expression of *MMP13* and Wnt target genes under Wnt stimulation. Using siRNA silencing, there was no effect of HIF1 $\alpha$ knockdown in COL2A and *Mmp13* in normoxia (Fig. 2*D*). Furthermore, HIF1 $\alpha$  knockdown has no effect on the nuclear translocation of EPAS1 (Fig. S2*B*).

Loss of HIF1 $\alpha$  promoted the Wnt-induced *Mmp13* expression and the transcriptional Wnt targets (Fig. 2*E*) and was confirmed by *Cre-lox* recombination (Fig. 2*F* and *G*). This was not observed when EPAS1 was knocked down (Fig. S2*C*).

To confirm that HIF1 $\alpha$  inhibits *Mmp13* expression by blocking transcription of Wnt targets, we stabilized HIF1 $\alpha$  in normoxic

chondrocytes upon von Hippel–Lindau tumor suppressor (*Vhl*) deletion using *Cre-lox* recombination. HIF1 $\alpha$  increase with VHL deletion was confirmed by Western blot analysis (Fig. 2*H*) and led to increased *Mmp13*, *Axin*, and WNT1–inducible-signaling pathway protein 1 (Wisp1) *expression* in normoxia (Fig. 2*I*). Moreover, stabilization of HIF1 $\alpha$  did not affect the transcription of the anabolic marker *COL2A1* (Fig. 2*I*). Thus, the regulation of anabolic markers is independent of the HIF1 $\alpha$  pathway. Because loss of VHL stabilizes both HIF1 and EPAS1, we used an overexpression of constitutive stabilized HIF1 in chondrocytes (18). We found that HIF1 $\alpha$  abolished *Mmp13* expression induced by Wnt (Fig. 2*I*). Thus, HIF1 $\alpha$  alone is able to inhibit Wnt-induced *Mmp13* expression.

HIF1 $\alpha$ - $\beta$ -Catenin Interaction Reduced TCF4 Binding to the *Mmp13* Regulatory Region. Upon Wnt pathway activation,  $\beta$ -catenin accumulates in the cytoplasm, translocates into the nucleus, binds to TCF transcription factor, and activates the transcription of target genes (19). Wnt activation increases the levels of catabolic markers under normoxia (16). We then investigated whether the translocation of  $\beta$ -catenin is reduced in hypoxia. Wnt3a promoted



**Fig. 2.** HIF1 $\alpha$  inhibits the transcription of Wnt targets in Wnt3a-induced chondrocytes. (A) Proteoglycan release in chondrocyte culture media (n = 9). qPCR analysis of relative gene expression in primary chondrocytes with 21% and 1% O<sub>2</sub> for: (B) anabolic marker (collagen 2A, *COL2A*) and catabolic marker (*Mmp13*) (n = 7); and (C) direct transcriptional targets of Wnt3a (*Axin* and *Wisp1*) (n = 14). (D) Expression of anabolic and catabolic genes (*COL2A* and *Mmp13*) with HIF1 $\alpha$  siRNA silencing (n = 6). (E) Direct transcriptional targets of Wnt3a (*Axin* and *Wisp1*) with HIF1 $\alpha$  siRNA silencing (n = 6). (E) Direct transcriptional targets of Wnt3a (*Axin* and *Wisp1*) with HIF1 $\alpha$  siRNA silencing (n = 6). (F) Catabolic marker *Mmp13* in HIF1 $\alpha$ -lacking chondrocytes (*Cre-lox* recombination in vitro) (n = 5). (G) Direct transcriptional targets of Wnt3a (*Axin* and *Wisp1*) with HIF1 $\alpha$ -lacking chondrocytes (*Cre-lox* recombination in vitro) (n = 5). (G) Direct transcriptional targets of Wnt3a (*Axin* and *Wisp1*) in HIF1 $\alpha$ -lacking chondrocytes (*Cre-lox* recombination in vitro) (n = 5). Data are mean  $\pm$  SEM. \*P < 0.05 compared with control, "P < 0.05. (H) Western blot analysis of HIF1 $\alpha$  expression in von Hippel-Lindau tumor suppressor protein (VHL)-lacking chondrocytes (with 21% O<sub>2</sub>), and quantification (n = 3). qPCR analysis of relative gene expression in VHL-lacking chondrocytes (*Cre-lox* recombination in vitro) for: (l) anabolic marker *COL2A*; catabolic marker *MMP13*; and direct transcriptional targets of Wnt (*Axin* and *Wisp1*) (with 21% O<sub>2</sub>). (J) Western blot analysis of stabilized HIF1 $\alpha$  expression using a tag antibody (HA) in primary chondrocytes and qPCR analysis of relative gene expression of *Mmp13*. Data are mean  $\pm$  SEM (n = 5 experiments); \*P < 0.05 compared with control; "P < 0.05.

 $\beta$ -catenin translocation into the nucleus and its protein expression regardless of O<sub>2</sub> level (Fig. 3 *A* and *B*). Thus, hypoxia reduced transcriptional activity of  $\beta$ -catenin independently of nuclear  $\beta$ -catenin translocation.

We investigated whether HIF1 $\alpha$  inhibits Wnt activity by direct interaction between HIF1 $\alpha$  and  $\beta$ -catenin. In hypoxia, HIF1 $\alpha$  and  $\beta$ -catenin colocalized in the nucleus of Wnt3a-induced chondrocytes (Fig. 4*A*), suggesting a possible interaction within nuclear complexes. Indeed, coimmunoprecipitation in nuclear extracts assay revealed HIF1 $\alpha$ - $\beta$ -catenin interaction complexes in hypoxia (Fig. 4*B*). However, EPAS1 failed to coimmunoprecipitate with  $\beta$ -catenin in normoxia and hypoxia (Fig. S2*D*). We next assessed the impact of HIF1 $\alpha$ - $\beta$ -catenin interaction on the transcriptional activity of TCF4- $\beta$ -catenin complexes. In chondrocyte stimulated by Wnt3a, the formation of TCF4- $\beta$ -catenin nuclear complexes was decreased in hypoxia compared with normoxia (Fig. 4*B*). These data suggest that under hypoxia,  $\beta$ -catenin may bind preferentially to HIF1 $\alpha$ rather than TCF4, thus reducing Wnt/ $\beta$ -catenin signaling.

MMP13 is regulated by both HIF and Wnt signaling (20–24). To better characterize the role of HIF1 $\alpha$ – $\beta$ -catenin interaction in inhibiting Wnt/ $\beta$ -catenin signaling and *Mmp13* expression under hypoxia, we assessed TCF4 binding to the *Mmp13* regulatory region by ChIP assay (Fig. 4C). We used a *Mmp13* regulatory 3' region downstream of coding area that includes Wnt responsive elements (WREs) (24) (Fig. 4C). TCF4 binding to WREs was decreased in hypoxia compared with normoxia (Fig. 4C). To confirm the role of HIF1 $\alpha$  in inhibiting TCF4 binding to WREs, we overexpressed the stabilized form of HIF1 $\alpha$  in normoxic chondrocytes and found that TCF4 binding to WREs was abolished



**Fig. 3.** Wnt3a promotes β-catenin translocation into the nucleus in hypoxia and normoxia. (A) Immunocytofluorescence staining of β-catenin in normoxic and hypoxic chondrocytes. Bar 100 μm (n = 4). Quantification of β-catenin translocation in chondrocytes: intensity of β-catenin signal into the nucleus of chondrocytes cultured in hypoxia and normoxia after Wnt3a stimulation (pixels) (n = 198-277). (B) Western blot analysis of β-catenin protein level in normoxic and hypoxic chondrocytes and quantification. Data are mean ± SEM. \*P < 0.05 compared with control (Ct).



**Fig. 4.** HIF1α binds β-catenin and inhibits TCF4 binding to the *MMP13* regulatory region. (A) Immunocytofluorescence staining of β-catenin and HIF1α in hypoxic chondrocytes (63×). (n = 3). (B) Coimmunoprecipitation of β-catenin in nuclear protein extracts. Western blot (WB) analysis of protein levels of HIF1α, TCF4, and β-catenin and quantification of HIF1α–β-catenin and TCF4–β-catenin complexes. (C) ChIP analysis of TCF4 binding to the *Mmp13* regulatory region. Sequence contains Wnt responsive elements. RNA Pol, RNA polymerase (positive control); IgG, mouse IgG (negative control). qPCR analysis of TCF4 binding to *Mmp13* regulatory regions (n = 3). \*P < 0.05 compared with control; \*P < 0.05. (D) Luciferase reporter assay in C3H10 cells. Data are ratio of firefly luciferase to control (Renilla) luciferase activity (n = 3). (E) qPCR analysis of relative gene expression (with HIF1α siRNA silencing) with 21% and 1% O<sub>2</sub> for *Mmp13* (n = 8) and direct transcriptional targets of Wnt (Axin and Wisp1) (n = 8). Data are mean  $\pm$  SEM. \*P < 0.05 compared with control; \*P < 0.05.

(Fig. 4*C*). Therefore, TCF4 binding to the *Mmp13* regulatory region was reduced by HIF1 $\alpha$ . We further investigated the transcriptional activity of *Mmp13* under hypoxia. The regulatory region was cloned downstream of a luciferase reporter gene and the plasmid was transfected in C3H10 cells. This assay revealed down-regulation of Wnt3a-induced luciferase activity when cultured under hypoxia (Fig. 4*D*) and confirmed the functional regulation of this sequence by hypoxia and Wnt3a.

**PKF118-310 Reduced OA Progression in**  $\Delta Hift \alpha^{chon}$  Mice. Given the sequestration of  $\beta$ -catenin by HIF1 $\alpha$ , we hypothesized that HIF1 $\alpha$  deletion may increase the TCF4– $\beta$ -catenin complex level, thereby leading to chondrocyte catabolism. To verify this hypothesis, we deleted HIF1 $\alpha$  in chondrocytes by siRNA silencing and blocked TCF4– $\beta$ -catenin complexes by using PKF118-310, which blocks the interaction between TCF4 and  $\beta$ -catenin. In normoxia, PKF118-310 totally suppressed Wnt-induced *Mmp13* and target gene expression (Fig. 4E). Loss of HIF1 $\alpha$  increased



**Fig. 5.** Loss of HIF1α increases TCF4–β-catenin complexes and cartilage lesions in OA mice with control or PKF118-310. (A) Safranin-O staining and OA score of *Hif1α<sup>THf1</sup>* and Δ*Hif1α<sup>chon</sup>* mouse knees after OA induction or control (Ct) at week 6 after treatment or not with PKF118-310. (Scale bar, 100 µm.) (B) MMP13 expression in *Hif1α<sup>THf1</sup>* and Δ*Hif1α<sup>chon</sup>* mice at week 6 and quantification. (Scale bar, 100 µm.) (C) TUNEL assay, quantification in *Hif1α<sup>fHf1</sup>* and Δ*Hif1α<sup>chon</sup>* mouse joints with OA or control at week 6. Data are mean ± SEM. \**P* < 0.05, \**P* < 0.05 compared with control Hif1α<sup>THf1</sup> (*n* = 8–11 animals per group).

the expression of *Mmp13* and that of the bona fide Wnt target genes *Axin* and *Wisp1*. Hence, the PKF118-310 addition inhibited the increased expression of the canonical Wnt targets *Mmp13*, *Axin*, and *Wisp1* induced by loss of HIF1 $\alpha$ . Thus, HIF1 $\alpha$ prevented Wnt from inducing *Mmp13* expression by blocking TCF4- $\beta$ -catenin complexes.

In OA mice, conditional loss of HIF1 $\alpha$  increased MMP13 expression and cartilage lesions (Fig. 5A). Increased TCF4- $\beta$ -catenin complex level may promote the phenotype observed in  $\Delta Hif1\alpha^{chon}$  mice. Articular injection of the  $\beta$ -catenin inhibitor PKF118-310 in  $\Delta Hif1\alpha^{chon}$  mice prevented cartilage lesions and reduced MMP13 expression compared with PBS treatment. Thus, PKF118-310 administration in  $\Delta Hif1\alpha^{chon}$  mice blocked the increased MMP13 expression and cartilage damage (Fig. 5A and B). Moreover, PKF118-310 resulted in a reduced number of apoptotic cells in OA mice (Fig. 5C). Loss of HIF1 $\alpha$  enhanced cartilage damage by increasing TCF4- $\beta$ -catenin complexes, which activated MMP13 expression.

# Discussion

Hypoxia is a characteristic of physiological articular cartilage (5, 25). We addressed the specific function of HIF1 $\alpha$  in chondrocytes and in OA in mice. With inducible conditional knockout of HIF1 $\alpha$  in mice, we showed that HIF1 $\alpha$  alleviated OA

development by down-regulating MMP13 through inhibition of  $\beta$ -catenin transcriptional activity in chondrocytes. The hypoxic avascular nature of the articular cartilage maintains the chondrocyte phenotype and homeostasis. Indeed, we observed that hypoxia and HIF1a were down-regulated in OA cartilage. Our findings are consistent with lower chondrocyte hypertrophy during hypoxia, which may contribute to the maintenance of cartilage homeostasis (26–28). Moreover, we show that HIF1 $\alpha$  is necessary to maintain a physiologic chondrocyte microenvironment and function. Consistent with our data, the inhibition of HIF1 $\alpha$  by 2-methoxyestradiol induced an OA phenotype in mice (5), which suggests that HIF1 $\alpha$  modulation is an important event that triggers chondrocyte differentiation in OA. With our conditional HIF1αknockout mouse model, we show that chondrogenic HIF1α directly maintains cartilage metabolism. Moreover, we show that HIF1 $\alpha$  is also a physiological antiapoptotic factor in articular cartilage as its loss enhanced chondrocyte apoptosis. Six weeks after HIF1α deletion, the increase of apoptotic chondrocyte did not induce cartilage lesion as chondrocyte apoptosis alone is not sufficient to induce cartilage lesion (15). Here, the increase in procatabolic enzymes in addition to increased apoptosis can exacerbate the cartilage lesion during OA. Finally the cartilage erosion observed in OA  $\Delta Hif1 \alpha^{chon}$  mice results from the double function of HIF1 $\alpha$ in articular cartilage as a Wnt inhibitor and an antiapoptotic protein. However, we cannot discard the possibility that Epas1 could be involved in cartilage erosion, because it was described as an inhibitor of survival and catabolic activator (22, 29).

Wnt/β-catenin signaling is one of the key pathways involved in OA (30). Its activation triggers the osteoarthritic differentiation of chondrocytes and OA in mice (16, 31). Thus, understanding the molecular regulators of Wnt signaling is of great therapeutic interest. Oxygen level is an important regulator of Wnt activity (9, 32). Indeed, high oxygen levels promote Wnt signaling and the differentiation of stem cells (33). Hypoxia inhibited the destruction of human cartilage explants by reducing MMP13 production in a HIF1 $\alpha$ -dependent manner (25). Because MMP13 is an important target of both HIF signaling and canonical Wnt pathways (20–24), we assessed whether HIF1a regulates MMP13 and Wnt signaling to prevent OA. Here we demonstrated that hypoxia down-regulated canonical Wnt signaling, thereby preventing chondrocyte catabolism. HIF1a deficiency exacerbated cartilage catabolism, thus HIF1 $\alpha$  is necessary to prevent OA development. We show that Wnt-induced MMP13 expression was promoted in  $\Delta Hif \alpha$ chondrocytes but was blunted by stabilized HIF1a in chondrocytes. We demonstrate that HIF1 $\alpha$  down-regulates the transcription of MMP13 driven by canonical Wnt signaling.

Given the critical role of β-catenin to drive Wnt signaling, understanding the role of HIF1 $\alpha$ - $\beta$ -catenin interaction may reveal novel mechanisms in regulating Wnt signaling. Because HIF1a inhibits TCF4– $\beta$ -catenin interaction and transcriptional activity (9), we investigated whether HIF1a modulates MMP13 through a Wnt/β-catenin pathway. Hypoxia promoted the translocation of  $\beta$ -catenin and binding to HIF1 $\alpha$  but not MMP13 expression. These results are consistent with transcriptional blockade of  $\beta$ -catenin activity despite the stabilization of  $\beta$ -catenin in the nucleus. Our study brings an unidentified mechanism into the protein regulation of HIF1 $\alpha$  and  $\beta$ -catenin interaction (9) and its impact on *Mmp13* transcription in chondrocytes. Under hypoxia, HIF1 $\alpha$ - $\beta$ -catenin complexes are preferentially formed, which results in a lower TCF4-\beta-catenin complex level and therefore markedly reduced TCF4 binding to the *Mmp13* regulatory region. Interestingly, when stabilized in normoxic chondrocytes, HIF1a blocked TCF4 binding to the Mmp13 regulatory region. Furthermore, the inhibition of TCF4-\beta-catenin complexes by PKF118-310 abolished Wnt3a-induced *MMP13* expression in  $\Delta$ HIF1 $\alpha$  chondrocytes. These results confirm the role of TCF4- $\beta$ -catenin complexes triggering MMP13 expression in  $\Delta$ HIF1 $\alpha$  chondrocytes. Thus, HIF1 $\alpha$ is a nuclear negative regulator of TCF4- $\beta$ -catenin complexes that inhibits the shift toward a catabolic phenotype in chondrocytes.

We further demonstrated that HIF1 $\alpha$  signaling protected against cartilage damage by blocking TCF4- $\beta$ -catenin. Indeed,

PKF118-310 administration reduced cartilage breakdown and the expression of MMP13 observed in  $\Delta Hif1\alpha^{chon}$  mice. Our findings support a procatabolic role of Wnt/β-catenin signaling in OA and bring new insights into the modulation of Wnt/β-catenin signaling by hypoxia. Furthermore, we provide evidence that HIF1α is a potent inhibitor of β-catenin and Wnt signaling and is required to block cartilage degradation. The use of HIF1α agonists might be a useful strategy for treating cartilage lesions in OA.

### Methods

Harvesting, Expansion, and Transfection of Primary Chondrocytes.  $Hif1a^{fllfl}$  and  $Vh^{fllfl}$  mice (The Jackson Laboratory) were used for chondrocyte cultures. Chondrocytes were harvested from 6-d-old mice and cultured with 10% (vol/vol) FBS. Recombination, transient overexpression, and plasmid (18) information are detailed in *SI Methods*.

**Real-Time PCR.** Real-time PCR involved use of SYBR green (Applied Biosystems) in five to eight independent experiments. Averaged Ct values were normalized to the averaged Ct value of *RpI13a*. Adjusted average Ct values were used to calculate relative expression versus control. Primer sequences are detailed in *SI Methods*.

**Immunocytochemistry.** Cells were cultured on cover glasses, fixed with 4% (wt/vol) formaldehyde. Cultures were saturated with 3% (wt/vol) BSA for 60 min, then incubated with the antibodies rabbit primary anti-HIF1 $\alpha$ , mouse primary anti- $\beta$ -catenin, and anti-EPAS1 (all Santa Cruz Biotechnology) for 1 h. Cultures were incubated for 60 min with Alexa dye 488-conjugated rabbit secondary antibody or Cy3 dye-conjugated mouse secondary antibody and observed with Axio Observer Z1 (Zeiss).

**Quantification of Proteoglycan Release, Western Blot Analysis, and Immunoprecipitation.** Proteoglycan release was measured in the supernatant by the colorimetric method. Proteins were extracted from whole cell lysates and nuclear extracts (n = 3 independent experiments). Immunoprecipitation, cloning, and reporter gene assay are described in *SI Methods*.

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Mice. To evaluate the expression of HIF1a during OA, we induced joint instability in 10-wk-old male C57BL6 mice (Janvier Labs) by DMM of the right knee and sham surgery at the left knee as described (34). Mice were killed at weeks 0 (n = 3), 4 (n = 5), and 6 (n = 5) after OA induction. fl/fl HIF1 $\alpha$ , R26R-LacZ, and Col2-Cre<sup>ERT</sup> mice were supplied by The Jackson Laboratory. At 9 wk of age, Col2-Cre<sup>ERT</sup>; fl/fl HIF1a, Col2-Cre<sup>ERT</sup>, and R26R-LacZ; and fl/fl HIF1a and R26R-LacZ mice were injected with tamoxifen (1 mg/10 g) (Sigma) daily for 5 d. Joint instability was induced in 10-wk-old male  $\Delta Hif1 \alpha^{chon}$  mice (n = 8-10 per group) and fl/fl HIF1 $\alpha$  littermates (n = 9-12 per group) in the right knee, with sham operation performed in the left knee.  $\varDelta Hif1\alpha^{chon}$  and fl/fl HIF1\alpha were injected intraarticularly once a week with PKF118-310 or PBS (n = 8 per group). Mice were killed 6 wk after surgery. This time point is suitable to quantify the expression of MMP13 in the remaining cartilage. To monitor hypoxia level in healthy and OA cartilage, we injected the hypoxia marker pimonidazole hydrochloride (Hypoxyprobe) in mice. Male FVB mice (The Jackson Laboratory) at 16 wk old were injected intraperitoneally with 0.6 mg pimonidazole/10 g weight and killed 17 h later. Mice were treated in accordance with the Guidelines for Animal Experimentation issued by the local committee (Lariboisière-Villemin no. CEEALV/2012-02-01, Paris).

**Histology.** Knees were fixed in 4% (wt/vol) PFA for 24 h at 4 °C and decalcified in Osteosoft; tissue was embedded in paraffin. Serial 5-µm-thick sagittal sections of medial femorotibial joints were collected at three depths at 70-µm intervals. Sections of tibias and femurs were stained with Safranin-O. The scoring method was used for tibias and femurs, with a total severity score ranging from 0 to 12 (35).

**Statistical Analysis.** Data are reported as mean  $\pm$  SEM. Statistical analyses involved ANOVA and the Mann–Whitney test (Statview, SAS Institute). *P* < 0.05 was the threshold of statistical significance.

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