



RESEARCH ARTICLE

2-Hydroxyisocaproic acid: A novel in vitro protease activity modulator with potential benefits for arthritis treatment

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ABSTRACT

Osteoarthritis affects approximately half a billion people worldwide and creates a significant economic burden, accounting for up to 2.5% of the national gross domestic product. Despite extensive research, a disease-modifying osteoarthritis drug remains unavailable. 2-hydroxyisocaproic acid is a physiological substance and the 2-hydroxy analogue of the essential amino acid leucine. This study investigates the potential of 2-hydroxyisocaproic acid to down-regulate key proteases involved in articular cartilage degradation. Specifically, we explore the use of 2-hydroxyisocaproic acid to inhibit the activity of matrix metalloproteinase 13 and a disintegrin and metalloproteinase with thrombospondin motifs 5, aiming to reduce the degradation of type II collagen and aggrecan, respectively, in osteoarthritis. Our findings demonstrate that 2-hydroxyisocaproic acid modulates and reduces the activity of both matrix metalloproteinase 13 and a disintegrin and metalloproteinase with thrombospondin motifs 5. Notably, 2-hydroxyisocaproic acid's reversible inhibition of these enzymes does not involve covalent bonding, positioning it as an enzyme modulator or down-regulator rather than a direct inhibitor.

Introduction

Arthritis manifests in over 100 distinct forms, among which osteoarthritis (OA) is predominant. This condition is characterized by structural impairments in hyaline articular cartilage (AC), compromised subchondral bone integrity, synovial tissue hypertrophy, enhanced vascularity, and instability of tendons and ligaments.^{1,2} The etiology of OA, however, remains not fully elucidated.³

Globally, OA was identified as the fourth leading cause of disability in 2020.⁴ By 2021, over 22% of individuals aged over 40 were suffering from knee OA, affecting an estimated 500 million people worldwide.⁵ In the United States alone, there were 54 million diagnosed cases and an additional 66 million self-reported cases in 2019.⁶ Beyond being a prevalent ailment second only to back pain, OA accounts for an annual financial impact of \$460 billion in medical expenses in the U.S.⁶ Beyond personal hardship, the socioeconomic implications in developed nations are profound, accounting for 1.0% to 2.5% of the gross domestic product.⁷

Emerging evidence suggests that the primary pathological events in OA involve changes in AC, leading to subsequent synovitis and subchondral bone degradation.⁸ It was not until the 1980s that the conceptual understanding of OA evolved from mere mechanical "wear and tear" degradation to a disease characterized by specific biochemical pathways causing AC damage.⁹⁻¹¹ OA is now recognized as a final common pathway influenced by multiple factors, most notably age, genetic predispositions, joint trauma, altered biomechanics, and obesity.¹²⁻¹⁵ This pathogenesis is likely triggered by trauma, leading to inflammation and the release of matrix-degrading enzymes,⁹ with altered biomechanical stress contributing to changes in chondrocyte metabolism, degradation of the specialized cartilage extra cellular matrix (ECM), and apoptosis of articular chondrocytes.¹⁶ Inflammation is now strongly linked to the pathogenesis of OA, with synovitis often arising as a secondary response

to cartilage damage, serving as a crucial link in the initiation and progression of the disease.¹²

Experimental studies, such as a single ex vivo impact to human cartilage, have shown cell death at the impact site, immediate release of inflammatory cytokines like interleukine-6 (IL-6) and tumor necrosis factor- α (TNF- α), radial progression of apoptosis, and cartilage degeneration extending to adjacent non-impacted areas.¹⁷ This disruption of chondrocytes' resting state may be viewed as an injury response that triggers developmental programs, leading to matrix remodeling, inappropriate hypertrophic maturation, and cartilage calcification.¹⁸

The initial loss of aggrecan is a critical early event in OA, beginning at the joint surface and extending to deeper zones, followed by collagen fibril degradation and mechanical tissue failure.¹⁹ Although proteoglycans are more remodeled under physiological conditions than collagens,²⁰ the sequence of degradation in the cartilage matrix components during OA development remains complex; collagen degradation typically follows the loss of aggrecan. Aggrecan loss can be reversed, but collagen degradation is irreversible, though cartilage cannot be repaired once collagen is lost.²¹ Eventually, a feedforward loop ensues as AC fragments, broken down by chondrocyte-derived proteinases, irritate the synovium, leading to secondary synovitis and further increases in proteases and inflammatory cytokines from synovial cells, thus perpetuating the disease cycle.²² OA is also associated with low-grade systemic and joint inflammation.^{18,23,24}

The metalloproteinase families, particularly a disintegrin and metalloproteinase with thrombospondin motifs (ADAMTSs) and matrix metalloproteinases (MMPs), are key mediators of cartilage destruction in OA.²²

The hallmark of OA pathology is extensive AC damage.²⁵ MMPs, a family of calcium-dependent and zinc-containing endoproteinases, vary in substrate specificity, cellular localization, activation,

and inducibility, belonging to the metzincin protease superfamily.¹ They possess broad substrate specificities, degrading not only ECM and basement membranes (BM) but also various bioactive non-matrix proteins, with 23 different types of MMPs encoded by 24 human genes.²⁶ MMPs also process and release cytokines, chemokines, insulin-receptor, and growth factors from their precursors or cryptic sites.^{27,28}

A principal collagenolytic enzyme, MMP-13 targets mature AC for degradation and is predominantly expressed in connective tissue.²⁹ It degrades type II collagen in cartilage, as well as proteoglycans, types IV and IX collagens, osteonectin, and perlecan in AC.³⁰ Clinical studies have associated high MMP-13 expression with AC destruction.³¹ MMP-13's role in early OA development is significant, being five to ten times more effective at degrading type II collagen than other collagenases.²² It plays a dual role in ECM destruction, degrading both collagen and the proteoglycan aggrecan.³²

The ADAMTS family of secreted zinc metalloproteinases comprises nineteen members known for their ability to bind and degrade extracellular cartilage matrix components.³⁰ ADAMTS-4 and -5 are the primary mediators of aggrecan cleavage in situ, playing significant roles in AC degradation.^{33,34}

LD-2-Hydroxy-isocaproic acid (HICA, also known as leucic acid), an organic acid with the chemical equation $C_6H_{12}O_3$ and a molecular weight of 132.2 g/mol, is an end product of leucine metabolism in human muscle and connective tissue, naturally occurring in mammalian organisms. During fasting, serum concentrations of HICA increase.³⁵

2-Hydroxyisocaproic acid, classified within the 2-hydroxycarboxylic acid group of amino acid metabolites, is part of the broader category of carboxylic acids, characterized by a carboxyl group (-COOH). This compound, among other carboxylic acids, has been reported to inhibit various MMPs,

particularly with derivatives of glutamine inhibiting MMP-8.³⁶ Some amino acid derivatives have shown in vitro effects on inhibiting connective tissue breakdown, and some molecules have been patented for treating degenerative joint diseases, though none are currently in clinical use.³⁷

In OA, the loss of aggrecan facilitates MMP-13 mediated degradation of exposed type II collagen. Both ADAMTS-5 and MMP-13 remain compelling therapeutic targets, provided that their side effects on physiological functions can be minimized.³⁸ Despite the absence of a disease-modifying osteoarthritis drug (DMOAD), the broad-spectrum, low-grade inhibition of MMPs could address the multifactorial nature of the disease more effectively than targeting a single enzyme, which might cause uncontrollable systemic damage.³⁹

Demonstrating efficacy in inhibiting a range of proteases, HICA has been effective against human MMP-2, -8, and -9, as well as neutrophil elastases and cathepsin G in vitro.^{40,41} This study aims to explore HICA's impact on key AC-degrading enzymes, offering a new potential therapeutic approach for OA.

Material and Methods

This experiment was carried out to demonstrate inhibition of recombinant human a disintegrin and metalloproteinase with thrombospondin motifs 5 (ADAMTS-5/Aggrecanase 2, #CC1034, Chemicon, International. Inc.) and human matrix metalloproteinase 13 (MMP-13, #30100802, BioTeZ, Berlin-Buch, Berlin, Germany) mediated break down of aggrecan (CC1890, Sigma-Aldrich, Darmstadt, Germany) by 10% rasemic D,L-2-hydroxy-isocaproic acid (HICA, Leios OÜ, Tallinn, Estonia) at pH 7.4. As positive inhibitor controls Ilomastat (Ilo, #CC1010, Millipore) and chlorhexidine (CHX, Corsodyl®, [2 mg/ml] (GlaxoSmithKline, Brøndby, Denmark) were used and 50 mM Tris-HCl-buffer (TNC): pH 7.5; 0.2 M NaCl; 1.0 mM $CaCl_2$) was used as neutral assay buffer solution in this study.⁴²

Briefly, MMP-13 was preincubated 1 hour at 37°C with p-aminophenylmercuric acetate (APMA, A9563, Sigma-Aldrich) to optimally activate (pro)MMP-13. As substrate, aggrecan was added and then ADAMTS-5 or with APMA activated MMP-13 as catalytic/peptolytic enzyme, TNC, as neutral control, and then 10% HICA, CHX or Ilo as protease/MMP-inhibitors were added. After that, they all were incubated together to find out the effects of different inhibitors on time function at a temperature of +37°C from 1 hour (h) to 2 days (d). Thus, the reactions were individually terminated by boiling with modified Laemmli's buffer. When the last incubation time was finished the samples were electrophoresed by 8% SDS-PAGE. After the electrophoresis the defragmentation of aggrecan was visualized using Silver Stain Pierce® (#24612, Thermo Fisher Scientific Inc., Vantaa, Finland). More details about the volumes of samples used in the experiment are described in the figure legends.

In experiment two MMP-13 activity was measured by catalytic activity assay as described previously.⁴³ In brief, MMP-13 was captured from assay buffer solution using MMP-13-specific monoclonal antibody-coated 96-well plates. The wells were

washed three times with PBS-T (phosphate buffered saline solution containing 0.05% [v/v] Tween-20 and incubated with 125-µl assay buffer (50 mM Tris-HCl, pH 7.6, 150 mM NaCl, 5 mM CaCl₂, 1 µM ZnCl₂ and 0,01% [v/v] Brij-35), to which 15 µl (50 µg/ml) modified pro-urokinase (UKcol) and 10 µl (6 mM stock) chromogenic substrate S-2444 was added. Color development was recorded by measurement of A₄₀₅ using a Titertek Multiskan 8-channel photometer. For measurement of total activity (already active plus latent MMP-13) in biological fluid, the immobilized MMP-13 was incubated With assay buffer containing 0.5 mM APMA for 2h, after which UKcol and S-2444 were added and activity was recorded.⁴³

Results

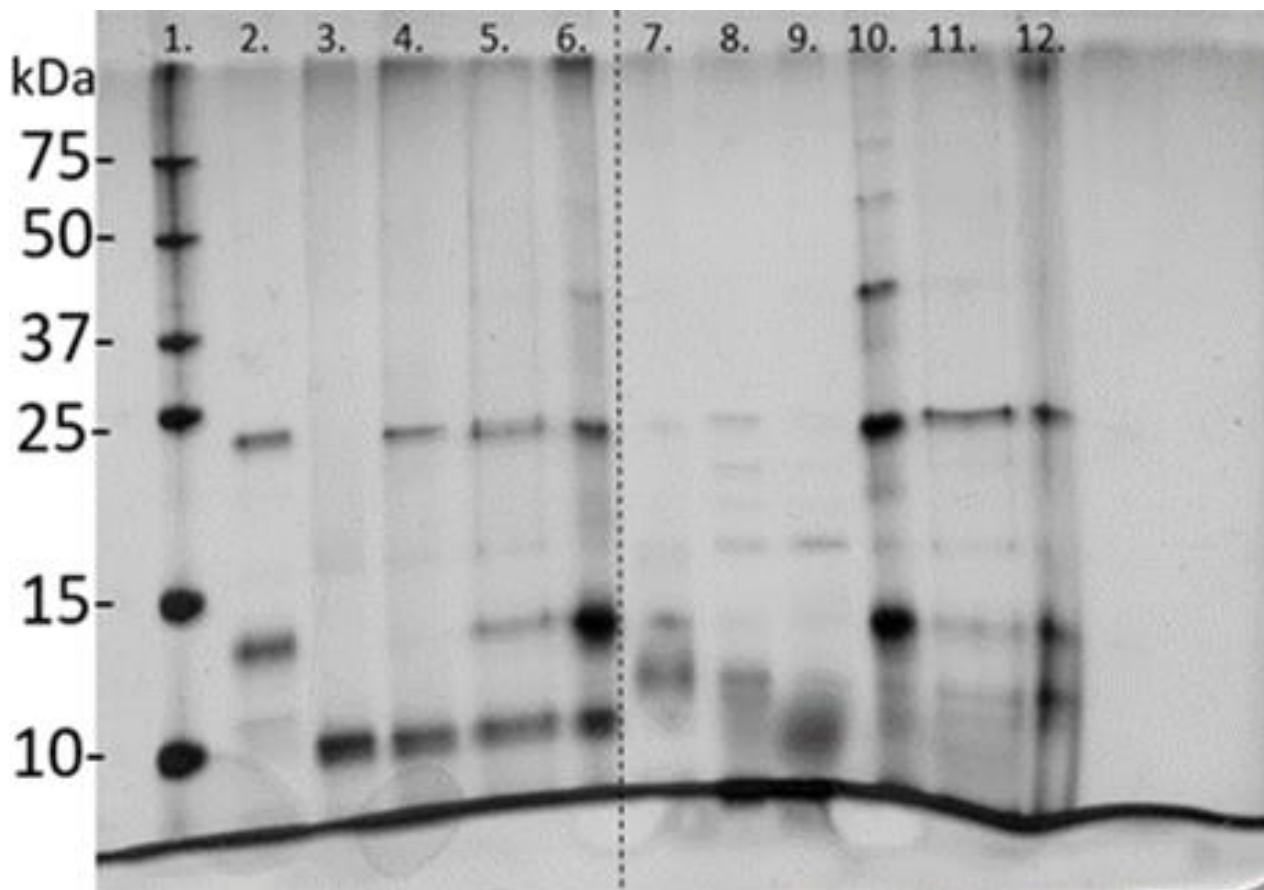
In the first experiment, we show that 10% D,L-2-hydroxy-isocaproic acid (HICA) inhibited the breakdown of aggrecan by both ADAMTS-5 and MMP-13 proteases (lanes 5 and 11, respectively). Figure 1 shows that HICA prevents break down of aggrecan. Therefore, both ADAMTS-5 and MMP-13 may be considered to be modulated enzyme activities by HICA rather than prevent enzyme activity like by chlorhexidine (CHX) in our study (Figure 1).

Figure 1. Illustrates the inhibition of recombinant human a disintegrin metalloproteinase with thrombospondin motifs 5 (ADAMTS-5) and human matrix metalloproteinase 13 (MMP-13) mediated break-down of aggrecan by 10% rasemic D,L-2-hydroxy-isocaproic acid (HICA). Molecular weight markers are indicated left. Figure 1 shows that 10% HICA inhibited the breakdown of aggrecan by both ADAMTS-5 and MMP-13 proteases (lines 5 and 11, respectively). Figure 1 shows that HICA prevents break down of aggrecan. Therefore, both ADAMTS-5 and MMP-13 may be considered to modulate enzyme activity rather than prevent enzyme activity like chlorhexidine in our study. Chlorhexidine (CHX). Ilomastat (Ilo)

Sample with concentrations in lines:

1. 0,25 µl molecular weight standards, Bio-Rad
2. 4,2 µl Aggrecan (0,5 µg/µl)
3. 4 µl ADAMTS-5 (0,1 µg/µl)
4. 4,2 µl Aggrecan + 4 µl ADAMTS-5 + 6 µl TNC-buffer, inc: 2d
5. 4,2 µl Aggrecan + 4 µl ADAMTS-5 + 6 µl 10% HICA, inc: 2d
6. 4,2 µl Aggrecan + 4 µl ADAMTS-5 + 6 µl CHX (2 mg/ml), inc: 2d
7. 1 µl MMP-13 (0,2 µg/µl) + 3 µl 2 mM APMA + 6 µl TNC
8. 4,2 µl Aggrecan + 1 µl MMP-13 (0,2 µg/µl) + 3 µl 2 mM APMA + 6 µl TNC, inc: 1h

- 9. 4,2 µl Aggrecan + 1 µl MMP-13 (0,2 µg/µl) + 3 µl 2 mM APMA + 6 µl TNC, inc: 2d
- 10. 4,2 µl Aggrecan + 1 µl MMP-13 (0,2 µg/µl) + 3 µl 2 mM APMA + 6 µl CXH, inc: 2d
- 11. 4,2 µl Aggrecan + 1 µl MMP-13 (0,2 µg/µl) + 3 µl 2 mM APMA + 6 µl 10% HICA, inc: 2d
- 12. 4,2 µl Aggrecan + 1 µl MMP-13 (0,2 µg/µl) + 3 µl 2 mM APMA + 6 µl Ilo. inc: 2d

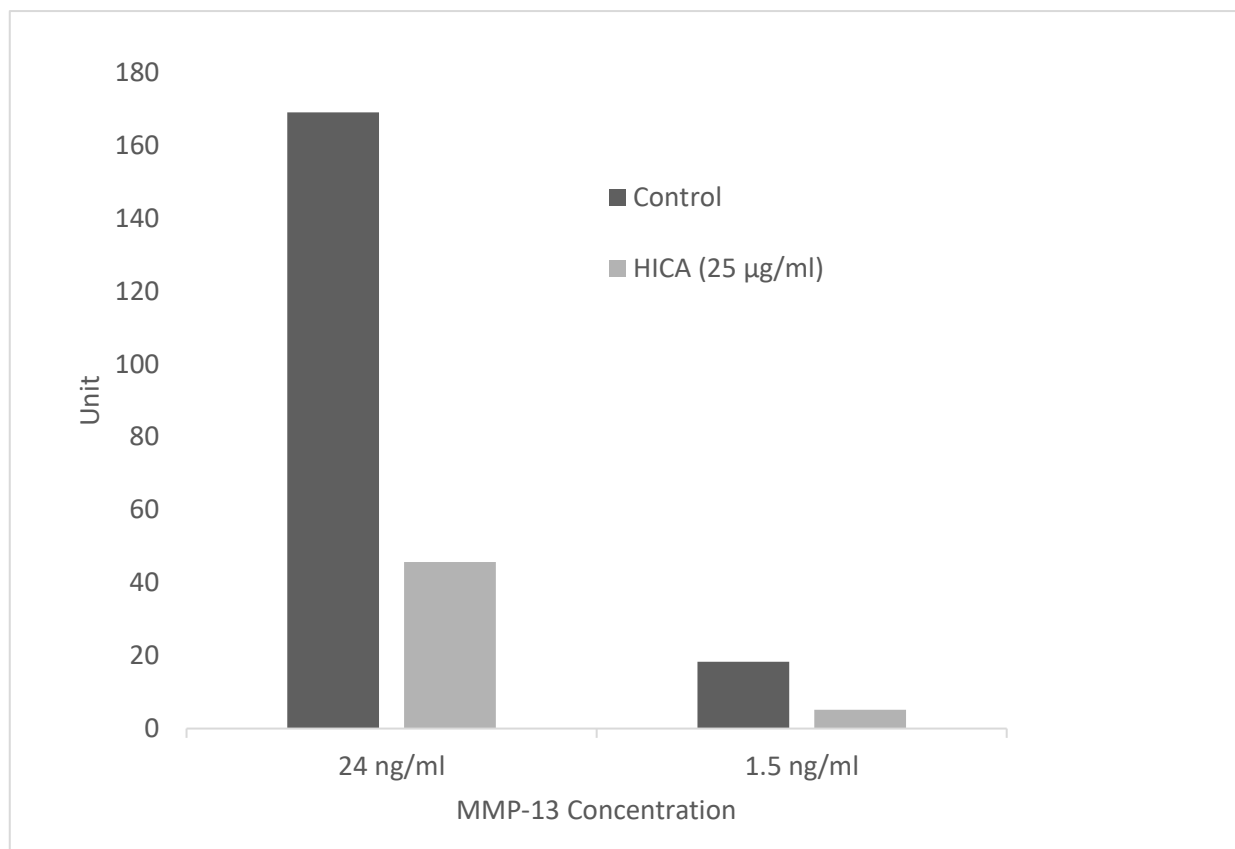


HICA's reversible inhibition of ADAMTS-5 and MMP-13 does not eventually involve covalent bonding and should thus be regarded as an enzyme modulator or down-regulator rather than a direct inhibitor.

In the second experiment, we demonstrated an inhibition of MMP-13 mediated break-down of

collagen by rasemic D,L-2-hydroxy-isocaproic acid (HICA), which is shown in Figure 2. HICA was found to exert effective inhibition of the activity of MMP-13 at a concentration level 25 µg/ml. At 24 ng/ml MMP-13 concentration there was 73% decreased activity noted and with 1.5 ng/ml concentration 72% decrease (Figure 2).

Figure 2. Inhibition of matrix metalloprotease 13 (MMP-13) activity by rasemic D,L-2-hydroxy-isocaproic acid (HICA) using modified pro-urokinase and chromogenic S2444 as substrate. MMP-13 activity was measured as described previously.⁴⁴ For a more detailed description, please refer to the Material and Methods section. HICA was found to effectively inhibit the activity of MMP-13 at a concentration of 25 µg/ml. At an MMP-13 concentration of 24 ng/ml, a 73% decrease in activity was noted, and at a concentration of 1.5 ng/ml, there was a 72% decrease.



Discussion

The aim of this study was to evaluate if HICA can inhibit key proteases involved in AC degradation. HICA was found to relatively inhibit or to down-regulate the human ADAMTS-5 aggrecanase when aggrecan served as the substrate. Similarly, human MMP-13 collagenase activity was also relatively inhibited, both when aggrecan and collagen were used as substrates.

Since the late 1970s, over 56 MMP inhibitors have been explored as clinical candidates, but only three have been clinically evaluated for cardiovascular indications: doxycycline for acute coronary syndrome and periodontal disease, Batimastat for coronary stent restenosis, and PG-116800 for preventing post-

ischemic left ventricular dilation.⁴⁵ To date, no MMP-13 inhibitor has reached clinical practice.⁴⁶ Since 2005, MMP inhibitors have been targeted for cancer (24 drugs), arthritis (27 drugs), and cardiovascular disease (10 drugs).⁴⁵

Inhibitors of matrix metalloproteinases MMPs could be effective in preventing the destruction of AC. Carboxylic acids, particularly tetrahydroisoquinoline-3-carboxylates, have shown promise in inhibiting MMP-8.⁴⁷ Nakai and colleagues from Scripps Research Institute identified potent MMP-13 inhibitors through high-throughput screening and activity-based protein profiling.⁴⁸ Adhikari and colleagues designed glutamine derivatives that initially showed potential in inhibiting MMPs.⁴⁹

Limited bioavailability and lack of enzyme selectivity have hindered the development of MMP inhibitors. Peptidomimetic compounds with a hydroxamic acid moiety were noted as very potent MMP inhibitors in the early phase of OA but still exhibited poor bioavailability and little specificity for targeted MMPs. Additionally, hydroxamic acid inhibitors are rapidly metabolized by the liver's first-pass metabolism and require frequent administration to maintain therapeutic plasma levels. More selective compounds are needed due to the overlap of function among individual enzymes and the complex inter-relationships involved in the expression, activation, and natural inhibition within the MMP family, which may necessitate multi-enzyme inhibition to ensure efficacy for targeted diseases.⁵⁰ Although PF152 seemed to be a very promising molecule that dose-dependently and selectively inhibited MMP-13 in experimental models, it was shown to cause nephrotoxicity by affecting the organic anion transporter 3.^{51,52}

Amino acid derivatives such as L-2-hydroxyisocaproate (HICA), a leucine analog, have been reported to influence protein synthesis similarly to L-leucine, exhibiting multiphasic responses that include inhibition of protein degradation at concentrations as low as 0.1 mM.⁵³ HICA has also been identified as an inhibitor of other proteinases⁵⁴ and is considered an anti-catabolic agent in clinical and experimental studies.⁵⁵⁻⁶⁴ HICA is classified as "anti-catabolite" and is widely used in the body building community.⁶⁴

A physiological substance, HICA has a plasma concentration in healthy adults of $0.25 \pm 0.02 \mu\text{M}$, and in circulation, it is not bound to plasma proteins.³⁵ Normal child serum concentration is $0.71 \pm 0.51 \mu\text{M}$ and range 0.02-2.04 (n=10).⁶⁵ It is found in human plasma, saliva, urine, amniotic fluid, and breast milk.⁶⁵⁻⁶⁹ Foods such as certain cheeses, sourdough, beer, radish, wines, and soy sauce also contain HICA.⁷⁰⁻⁷⁹

Effectively absorbed through active transport in the human bowel, HICA can bypass the liver's first-pass

metabolism.⁸⁰⁻⁸² It can also be assumed that HICA is also passively absorbed from alimentary canal by diffusion.^{62,63,83} Thus, HICA does reach the target tissue, the systemic circulation, when administered orally.^{84,85} Topically, HICA is non-toxic even at high doses and HICA is not cytotoxic nor genotoxic at concentrations $<10 \text{ mg/ml}$, regardless of the exposure time, although it was cytostatic at all tested concentrations. HICA retained about 70% of the osteogenic differentiation potential at 1 mg/ml.^{86,87}

Aggrecan, the predominant proteoglycan in cartilage, is composed of chondroitin sulphate and keratan sulphate binding to the linear core protein hyaluronic acid backbone. It provides AC structural rigidity, compressibility, and collagen elasticity.^{88, 89} Aggrecan gives a low coefficient of friction on the joint surfaces and gives the cartilage protective resiliency.²²

Two major families within the metalloproteinase class, namely a disintegrin and metalloproteinase with thrombospondin motifs (ADAMTSs) and the matrix metalloproteinases (MMPs), are principal agents of cartilage destruction in arthritic conditions.²² Specifically, ADAMTS-4 and -5 are pivotal in the cleavage of aggrecan in situ, significantly contributing to cartilage degradation.^{33,34} In healthy joints, distinct locations within the cartilage are mapped by MMP and aggrecanase neoepitopes, indicating that these proteolytic enzyme groups operate at different sites during the normal turnover of aggrecan.¹⁹

The degradation of AC in arthritic diseases is primarily mediated by MMPs and ADAMTSs, with MMP-13 being a major collagenolytic enzyme in this process (Vincenti and Brinckerhoff 2002). It is critical for the progression of OA and is a prime target for pharmacological intervention.²⁵

Population growth, especially among the elderly, and rising obesity rates do not entirely account for the sharp increase in total knee replacements observed in recent decades. It is reasonable to assume that additional factors are at play. A

notably high increase in knee replacements among younger individuals is likely linked to an increase in adolescent knee injuries and broader criteria for undergoing this procedure.⁹⁰ Globally, joint replacement surgeries are expanding at an annual rate of 10%, with 95% performed on patients with OA.⁹¹ Addressing therapeutic targets at various OA stages is critically needed.⁴

Despite successful anterior cruciate ligament (ACL) repairs, which involve reconstruction of torn ligaments, 45%-50% of these patients will eventually develop OA.⁹²⁻⁹⁴ Studies comparing OA prevalence between reconstructed and healthy, operated knees show a higher incidence of OA.⁹⁵ This suggests that ACL reconstruction does not prevent OA, with a threefold increase in OA prevalence observed in knees post-reconstruction compared to their healthy counterparts.

Bluteau et al.⁹⁶ observed a significant surge in MMP-13 and MMP-3 expression following ACL surgery, with no change in the expression of tissue inhibitors of MMPs (TIMP). The expression of MMP-13 genes rose sharply post-surgery and remained elevated. Similarly, a marked increase in MMP-13 gene expression was noted immediately after ACL transection, which continued at high levels. Corresponding increases in MMP-13 mRNA and protein were observed in the synovium and meniscus. The joint cavity releases substantial amounts of MMPs into the joint fluid, potentially disrupting the balance between tissue synthesis and degradation, thus impeding ACL healing and exacerbating osteoarthritis.⁹⁷

The inhibition of ADAMTS-5 may safeguard against cartilage damage and aggrecan loss following OA induction via surgical destabilization of the medial meniscus.⁹⁸ It is noteworthy that blocking ADAMTS-5 activity might also reduce pain, according to preclinical studies.⁸⁸

Conclusion

Proteases play a crucial role in normal physiological processes, making their complete inhibition unfeasible. Despite the incomplete understanding of the arthritic disease and its destructive progression, it is essential to modulate excessive protease activity to restore physiological balance. Although significant efforts have been made to develop effective therapies or disease-modifying drugs for osteoarthritis and its progression, success has been limited. This study suggests that D,L-2-hydroxy-isocaproic acid, with its ability to modulate enzyme activity and inhibit key proteinases, may offer therapeutic benefits across all stages of OA. The aim of this study was to evaluate the potential of HICA to inhibit key proteases involved in AC degradation. HICA was found to significantly inhibit or down-regulate human ADAMTS-5 aggrecanase activity when aggrecan was used as the substrate. Similarly, human MMP-13 collagenase activity was also significantly inhibited.

Disclosure of potential conflicts of interest:

The authors are named as inventors on patents FI129515B and U.S. Patent No. 11,793,779 both of which were filed by Salarusta Ltd, these patents are related to this work and the patents cover the use of a substance on a treatment of arthritis. The authors have an ownership interest in Salarusta Ltd Oy. The authors have no other competing interests such as relevant affiliations or financial involvement with any organization or entity with a financial interest in or financial conflict with the subject matter or materials discussed in the manuscript. This includes employment, consultancies, honoraria, stock ownership or options, expert testimony, grants, or patents received or pending, or royalties.

The data presented herein was previously disclosed in part in the aforementioned Patent FI12951B and U.S. Patent No. 11,793,779. In this study, we further explore the findings initially disclosed in the Patents, where we presented the preliminary data on 2-hydroxyisocaproic acid's effects on various proteinases. This paper expands upon these findings by more practical approach to tackle proteinases relevant to arthritis especially osteoarthritis.

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